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2 Ocean biogeochemistry in the Canadian Earth System Model version 5.0.3: CanESM5 and
3 CanESM5-CanOE

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5 James R. Christian^{1,2}, Kenneth L. Denman^{2,3}, Hakase Hayashida^{3,4}, Amber M. Holdsworth¹,
6 Warren G. Lee², Olivier G.J. Riche^{3,5}, Andrew E. Shao^{2,3}, Nadja Steiner^{1,2}, and Neil C. Swart²

7

8 1 Fisheries and Oceans Canada, Sidney, BC, Canada

9 2 Canadian Centre for Climate Modelling and Analysis, Victoria, BC, Canada

10 3 School of Earth and Ocean Sciences, University of Victoria, Victoria, BC, Canada

11 4 now at the Institute for Marine and Antarctic Studies, University of Tasmania, Hobart,
12 Tasmania, Australia

13 5 now at Fisheries and Oceans Canada, Mont Joli, Québec, Canada

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16 *Correspondence to:* James Christian (jim.christian@ec.gc.ca)

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18



19 **Abstract.** The ocean biogeochemistry components of the Canadian Earth System Model v. 5 are
20 presented and compared to observations and other models. CanESM5 employs the same
21 biogeochemistry module as CanESM2 whereas CanESM5-CanOE (“Canadian Ocean Ecosystem
22 model”) is a new, more complex biogeochemistry module developed for CMIP6, with multiple
23 food chains, flexible phytoplankton elemental ratios, and a prognostic iron cycle. This new
24 model is described in detail and the outputs compared to CanESM5 and CanESM2, as well as to
25 observations and other CMIP6 models. Both CanESM5 models show gains in skill relative to
26 CanESM2, which are attributed primarily to improvements in ocean circulation. CanESM5-
27 CanOE shows improved skill relative to CanESM5 in some areas. CanESM5-CanOE includes a
28 prognostic iron cycle, and maintains high nutrient / low chlorophyll conditions in the expected
29 regions (in CanESM2 and CanESM5, iron limitation is specified as a temporally static ‘mask’).
30 Surface nitrate concentrations are biased low in the subarctic Pacific and equatorial Pacific, and
31 high in the Southern Ocean. Export production in CanESM5-CanOE is among the lowest for
32 CMIP6 models; in CanESM5 it is among the highest, but shows the most rapid decline after
33 about 1980. CanESM5-CanOE has relatively low concentrations of zooplankton and detritus
34 relative to phytoplankton, and a high and relatively constant living phytoplankton fraction of
35 total particulate organic matter. In most regions, large and small phytoplankton show decoupled
36 seasonal cycles with greater abundance of large phytoplankton in the productive seasons.
37 Cumulative ocean uptake of anthropogenic carbon dioxide through 2014 is lower in both
38 CanESM5 models than in observation-based estimates or the model ensemble mean, and is lower
39 in CanESM5-CanOE (122 PgC) than in CanESM5 (132 PgC).

40



41 **1. Introduction**

42

43 The Canadian Centre for Climate Modelling and Analysis has been developing coupled models
44 with an interactive carbon cycle for more than a decade (Arora et al., 2009; 2011; Christian et al.,
45 2010). CanESM5 (Swart et al., 2019a) is an updated version of CanESM2 (Arora et al., 2011),
46 with an entirely new ocean and an atmosphere with the same T63 horizontal resolution and
47 important improvements in atmospheric physics. The CanESM5 ocean is based on the Nucleus
48 for European Modelling of the Ocean (NEMO) system version 3.4. The ocean biogeochemistry
49 modules were developed in-house, although parameterizations for some processes were adapted
50 from the native PISCES biogeochemistry model (Aumont et al., 2015). CanESM5 uses the same
51 biogeochemistry model as CanESM1 and CanESM2, the Canadian Model of Ocean Carbon
52 (CMOC; Zahariev et al., 2008), adapted for use within NEMO. An additional model was
53 developed for CMIP6, called the Canadian Ocean Ecosystem model (CanOE). The biological
54 components of CanOE are of substantially greater complexity, including multiple food chains,
55 flexible phytoplankton elemental ratios, and a prognostic iron (Fe) cycle. Carbon chemistry, gas
56 exchange and solubility of carbon dioxide (CO₂) and oxygen are identical between the two and
57 follow the protocols specified by the Ocean Model Intercomparison Project - Biogeochemistry
58 (OMIP-BGC) (Orr et al., 2017). The two coupled models are known as CanESM5 and
59 CanESM5-CanOE, respectively. There are no feedbacks between biology and the physical ocean
60 model, so the physical climate of CanESM5 and CanESM5-CanOE is identical in experiments
61 with prescribed atmospheric CO₂ concentration.

62

63 The reasons for developing both models are, firstly, to evaluate the effect of changes in ocean



64 circulation between CanESM2 and CanESM5 on ocean biogeochemistry by running the new
65 climate model with the same ocean biogeochemistry, and secondly because CanOE is
66 substantially more expensive computationally (19 tracers vs 7). Having CMOC as an option
67 allowed us to run many CMIP6 experiments with CanESM5 only, as ocean biogeochemistry is
68 not central to their purpose. Many additional tracers requested by OMIP-BGC including abiotic
69 and natural dissolved inorganic carbon (DIC), DI^{14}C , CFCs and SF_6 (see Orr et al., 2017) were
70 run only in CanESM5, since adding these tracers on top of the larger set of biological
71 components in CanOE would have been prohibitively expensive. The CMIP6 experiments
72 published for CanESM5-CanOE are listed in Supplementary Table S1.

73

74 CMOC is a nutrient-phytoplankton-zooplankton-detritus (NPZD) model with highly
75 parameterized representations of phytoplankton Fe limitation, dinitrogen (N_2) fixation and
76 denitrification, and calcification and calcite dissolution (Zahariev et al., 2008). In CanESM1 and
77 CanESM2, CMOC did not include oxygen. In CanESM5, CMOC now includes oxygen as a
78 purely 'downstream' tracer that does not affect other biogeochemical processes, whereas in
79 CanOE denitrification is prognostic and dependent on the concentration of oxygen. Among the
80 less satisfactory aspects of CMOC biogeochemistry are, firstly, that Fe limitation is specified as a
81 static 'mask' that does not change with climate (being calculated from the present-day
82 climatological distribution of nitrate), and secondly, that denitrification is parameterized so that
83 nitrogen (N) is conserved within each vertical column, i.e., collocated with N_2 fixation in tropical
84 and subtropical open-ocean regions (Zahariev et al., 2008; Riche and Christian, 2018). This latter
85 simplification produced excessive accumulations of nitrate in Eastern Boundary Current regions
86 where most denitrification actually occurs. CMOC also has a tendency to produce rather stark



87 extremes of high and low primary and export production (Zahariev et al., 2008), a well-known
88 problem of NPZD models (Armstrong, 1994; Friedrichs et al., 2007). Our intent in developing
89 CanOE was to alleviate, or at least reduce, these biases, by including multiple food chains, a
90 prognostic Fe cycle, and prognostic denitrification. Dinitrogen fixation is still parameterized, but
91 the CanOE parameterization includes Fe limitation, whereas in CMOC N₂ fixation tends to grow
92 without bound in a warming ocean as there is no P or Fe limitation (Riche and Christian, 2018).
93 Calcification is represented by a prognostic detrital calcite pool with its own sinking rate (distinct
94 from that of organic detritus), and calcite burial depends on the saturation state. In CMOC
95 calcification is parameterized by a temperature dependent “rain ratio” and 100% burial of calcite
96 that reaches the seafloor is assumed.

97

98 In this paper we present a detailed model description for CanOE and an evaluation of both
99 CanESM5 and CanESM5-CanOE relative to observational data products and other available
100 models. CMOC has been well described previously (Zahariev et al., 2008) and the details are not
101 reiterated here. In some cases, CanESM2 results are also shown to illustrate which differences in
102 the model solutions arise largely from the evolution of the physical ocean model, and which are
103 specifically associated with different representations of biogeochemistry.

104

105 **2. Model Description**

106

107 The NEMO system is a publicly available archive of codes based on the OPA (Océan
108 PARallelisé) ocean model (Madec and Imbard, 1996; Guilyardi and Madec, 1997). It comes with
109 two options for biogeochemistry: PISCES (Pelagic Interactions Scheme for Carbon and



110 Ecosystem Studies) and LOBSTER (LODyC Ocean Biogeochemical System for Ecosystem and
111 Resources). CanOE and the NEMO implementation of CMOC are built around the basic code
112 structure of PISCES within the Tracers in Ocean Paradigm (TOP) module, using NEMO v3.4.1,
113 but have also been implemented in NEMO 3.6 for regional downscaling applications
114 (Holdsworth et al., 2021).

115

116 The biology, carbon chemistry, gas exchange and light attenuation components have all been
117 modified to various degrees. In a few cases PISCES parameterizations, or slightly modified
118 versions thereof, were adopted. CanOE uses PISCES three-band light attenuation while NEMO-
119 CMOC uses broadband attenuation of photosynthetically active radiation (PAR) for consistency
120 with the published version of CMOC. Carbon chemistry was modified to be consistent with the
121 Best Practices Guide (Dickson et al., 2007) and the OMIP-BGC data request (Orr et al., 2017).
122 All calculations are done on the total scale and the recommended formulae for the equilibrium
123 constants are employed. The PISCES conventions for convergence of carbon chemistry
124 calculations were retained, the greater number of iterations in the surface layer offering greater
125 accuracy in calculating $p\text{CO}_2$ and gas exchange. In subsurface layers (where the only function of
126 the carbon chemistry is to calculate burial of calcite in the sediments) the number of iterations is
127 fixed at five. CanESM5 uses the same carbon chemistry but does not solve the carbon chemistry
128 equations in the subsurface layers.

129

130 The CanOE biology model is a substantially new model based on the cellular regulation model
131 of Geider et al. (1998). There are two phytoplankton functional groups, and each group has four
132 state variables: C, N, Fe and chlorophyll. Photosynthesis is decoupled from cell production and



133 photosynthetic rate is a function of the cell's internal N and Fe quotas. Each functional group has
134 a specified minimum and maximum N quota and Fe quota, and nutrient uptake ceases when the
135 maximal cell quota is reached. Chlorophyll synthesis is a function of N uptake and increases at
136 low irradiance. Model parameters and their values are listed in Table 1. A schematic of the
137 model is shown in Figure 1.

138

139 **2.1 Photosynthesis and Phytoplankton Growth**

140

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

144

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

$$146 \quad T_f = \exp\left(-\frac{E_{ap}}{R} \left(\frac{1}{T} - \frac{1}{T_{ref}}\right)\right) \quad (1)$$

147 where E_{ap} is an enzyme activation energy that corresponds approximately to that of RuBisCo (cf.
148 Raven and Geider 1988), R is the gas constant ($8.314 \text{ J mol}^{-1} \text{ K}^{-1}$), and temperature T and
149 reference temperature T_{ref} are in Kelvin. Maximal rates of nutrient (either N or Fe, but
150 generically referred to here with the superscript X) uptake are given by

$$151 \quad V_{max}^X = V_{ref}^X T_f \left(\frac{Q_{max}^X - Q^X}{Q_{max}^X - Q_{min}^X}\right)^{0.05} \quad (2)$$

152 where V_{max}^X is the maximal uptake rate in mg of nutrient X per mg of cell C, X can represent N
153 or Fe, Q is the nutrient cell quota and Q_{min} and Q_{max} its minimum and maximum values, and V_{ref}^X

154

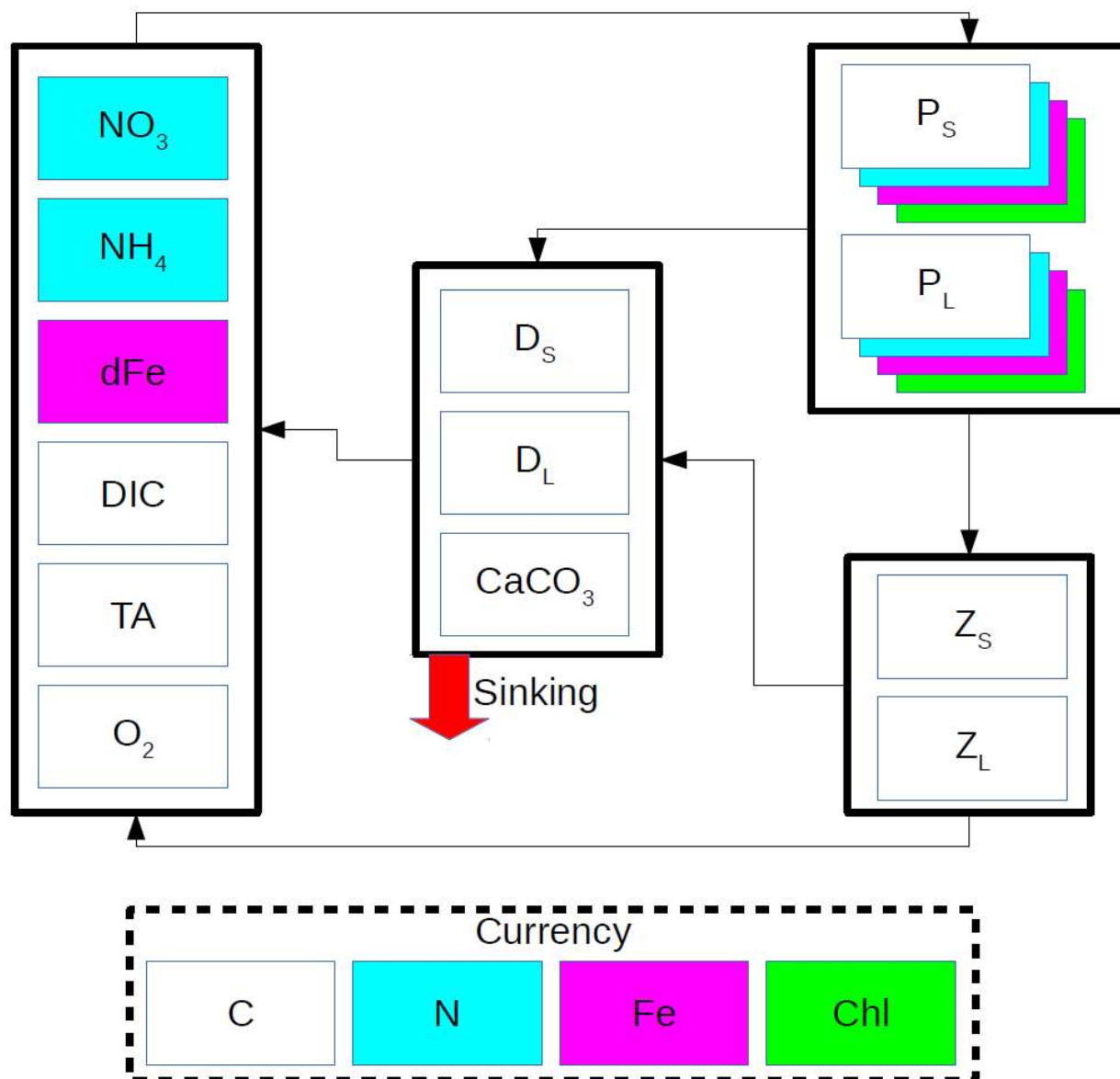


Figure 1 - Schematic of the CanOE biology model. Model currencies including chlorophyll (Chl) are indicated by coloured boxes except oxygen (O_2) and carbonate (CaCO_3). Arrows indicate flows of carbon (C), nitrogen (N) and iron (Fe) between compartments containing small (S) and large (L) phytoplankton (P), zooplankton (Z), and detritus (D) components; counterflows of oxygen are not shown.



Table 1 – Ecosystem model parameters.

Symbol	Description	Unit	
T_{ref}	Reference temperature	K	298.15
E_{ap}	Activation energy for photosynthesis	kJ mol^{-1}	37.4
$Q_{\text{mins}}^{\text{N}}$	Small phytoplankton minimum N quota	g N g C^{-1}	0.04
$Q_{\text{maxs}}^{\text{N}}$	Small phytoplankton maximum N quota	g N g C^{-1}	0.172
$Q_{\text{minl}}^{\text{N}}$	Large phytoplankton minimum N quota	g N g C^{-1}	0.04
$Q_{\text{maxl}}^{\text{N}}$	Large phytoplankton maximum N quota	g N g C^{-1}	0.172
$Q_{\text{mins}}^{\text{Fe}}$	Small phytoplankton minimum Fe quota	$\mu\text{g Fe g C}^{-1}$	4.65
$Q_{\text{maxs}}^{\text{Fe}}$	Small phytoplankton maximum Fe quota	$\mu\text{g Fe g C}^{-1}$	93.
$Q_{\text{minl}}^{\text{Fe}}$	Large phytoplankton minimum Fe quota	$\mu\text{g Fe g C}^{-1}$	6.5
$Q_{\text{maxl}}^{\text{Fe}}$	Large phytoplankton maximum Fe quota	$\mu\text{g Fe g C}^{-1}$	70.
$V_{\text{ref}}^{\text{N}}$	Reference rate of N uptake	$\text{g N g C}^{-1} \text{d}^{-1}$	0.6
$V_{\text{ref}}^{\text{Fe}}$	Reference rate of Fe uptake	$\mu\text{g Fe g C}^{-1} \text{d}^{-1}$	79.
$P_{\text{ref}}^{\text{C}}$	Reference rate of photosynthesis	$\text{g C g C}^{-1} \text{d}^{-1}$	3
k_{XU}	Rate coefficient for exudation	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	$\text{g C g CHL}^{-1} \text{h}^{-1} (\mu\text{mol m}^{-2} \text{s}^{-1})^{-1}$	1.08
$\Theta_{\text{max}}^{\text{N}}$	Maximum chlorophyll-nitrogen ratio	g g^{-1}	0.18
K_{NiS}	Half-saturation for small phytoplankton nitrate uptake	$\text{mmol}^{-1} \text{m}^3$	0.1
K_{NaS}	Half-saturation for small phytoplankton ammonium uptake	$\text{mmol}^{-1} \text{m}^3$	0.05
K_{FeS}	Half-saturation for small phytoplankton iron uptake	$\text{nmol}^{-1} \text{m}^3$	100
K_{NiL}	Half-saturation for large phytoplankton nitrate uptake	$\text{mmol}^{-1} \text{m}^3$	1.0
K_{NaL}	Half-saturation for large phytoplankton ammonium uptake	$\text{mmol}^{-1} \text{m}^3$	0.05
K_{FeL}	Half-saturation for large phytoplankton iron uptake	$\text{nmol}^{-1} \text{m}^3$	200
$m_{1\text{S}}$	Small phytoplankton/zooplankton mortality rate (linear)	d^{-1}	0.05
$m_{2\text{S}}$	Small phytoplankton/zooplankton mortality coefficient	$(\text{mmol C m}^{-3})^{-1} \text{d}^{-1}$	0.06
$m_{1\text{L}}$	Large phytoplankton/zooplankton mortality rate (linear)	d^{-1}	0.1



m_{2L}	Large phytoplankton/zooplankton mortality coefficient	$(\text{mmol C m}^{-3})^{-1} \text{ d}^{-1}$	0.06
X_{minp}	Minimum phytoplankton concentration for linear mortality	mmol C m^{-3}	0.01
a_L	Large zooplankton grazing parameter	$(\text{mmol C m}^{-3})^{-1}$	0.25
G_{L0}	Large zooplankton maximum grazing rate	d^{-1}	0.85
a_S	Small zooplankton grazing parameter	$(\text{mmol C m}^{-3})^{-1}$	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
E_{ar}	Activation energy for detritus remineralization	kJ mol^{-1}	54.0
w_s	Small detritus sinking speed	m d^{-1}	2.
w_l	Large detritus sinking speed	m d^{-1}	30.
w_{Ca}	CaCO_3 sinking speed	m d^{-1}	20.
P_{Ca}	CaCO_3 production as fraction of mortality	$\text{mol CaCO}_3 \text{ molC}^{-1}$	0.05
k_{Ca}	CaCO_3 dissolution rate	d^{-1}	0.0074
$S_{\text{Fe}1}$	Dissolved iron scavenging loss rate ($\text{Fe} \leq L_{\text{Fe}}$)	d^{-1}	0.001
$S_{\text{Fe}2}$	Dissolved iron scavenging loss rate ($\text{Fe} > L_{\text{Fe}}$)	d^{-1}	2.5
L_{Fe}	Ligand concentration	nmol Fe m^{-3}	600.
P_{Fe}	POC-dependence parameter for Fe scavenging	$(\text{mmolC m}^{-3})^{-1}$	0.66
$K_{\text{NH}4\text{ox}}$	Nitrification rate in darkness	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	$\text{mmol m}^{-3} \text{ d}^{-1}$	0.0225
a	Initial slope for irradiance-dependence of N_2 fixation	$(\text{W m}^{-2})^{-1}$	0.02
K_{Fe}	Half-saturation for Fe dependence of N_2 fixation	nmol m^{-3}	100.
$K_{\text{NO}3}$	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



158 is a (specified) basal rate at $T=T_{ref}$ and $Q=Q_{min}$. These maximum rates are then reduced according
 159 to the ambient nutrient concentration, i.e.

$$160 \quad V^N = V_{max}^N (L_{NH4} + (1 - L_{NH4})L_{NO3}) \quad (3a)$$

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

$$163 \quad V^{Fe} = V_{max}^{Fe} \left(\frac{Fe}{K_{FeX} + Fe} \right) \quad (3b)$$

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

$$166 \quad P_{max}^C = P_{ref}^C T_f \min \left\{ \frac{Q^N - Q_{min}^N}{Q_{max}^N - Q_{min}^N} \cdot \frac{Q^{Fe} - Q_{min}^{Fe}}{Q_{max}^{Fe} - Q_{min}^{Fe}} \right\} \quad (4)$$

167 where P_{ref}^C is the rate at the reference temperature T_{ref} under nutrient-replete conditions
 168 ($Q=Q_{max}$). The light-limited growth rate is then given by

$$169 \quad P_{phot}^C = P_{max}^C \left(1 - e^{-\alpha_{chl} E \theta_C / P_{max}^C} \right) \quad (5)$$

170 where θ_C is the chlorophyll-to-carbon ratio. The rate of chlorophyll synthesis is

$$171 \quad \rho_{chl} = \theta_{max}^N \frac{P_{phot}^C}{E \alpha_{chl} \theta} \quad (6)$$

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

$$174 \quad \frac{dC_p}{dt} = (P_{phot}^C - \zeta V_N) C_p - (G + C_{XS}) - m_1 C_p - m_2 C_p^2 - k_{XU} C_{INTR} \quad (7)$$

175 where ζ is the respiratory cost of biosynthesis, G is the grazing rate (equation 12), C_{XS} is the
 176 excess (above the ratio in grazer biomass) carbon in grazing losses, m_1 and m_2 are coefficients
 177 for linear and quadratic nonspecific mortality terms, C_{INTR} is the concentration of intracellular



178 carbohydrate carbon in excess of biosynthetic requirements, and k_{XU} is a rate coefficient for its
179 exudation to the environment. The nonspecific mortality terms are set to 0 below 0.01 mmol C
180 m^{-3} , to prevent biomass from being driven to excessively low levels in the high latitudes in
181 winter (Hayashida, 2018). The full equation for phytoplankton N, Fe and chlorophyll are

$$182 \quad \frac{dN_p}{dt} = \frac{V^N}{Q_N} - (G + m_1 C_p + m_2 C_p^2) R_{NC} - N_{XS} \quad (8)$$

$$183 \quad \frac{dFe_p}{dt} = \frac{V^{Fe}}{Q_{Fe}} - (G + m_1 C_p + m_2 C_p^2) R_{FeC} - Fe_{XS} \quad (9)$$

$$184 \quad \frac{dM}{dt} = \frac{\rho_{chl} V^N}{\theta_C} M - (G + m_1 C_p + m_2 C_p^2) \theta_C - k_{dgr} M \quad (10)$$

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

189

190 **2.2 Grazing and Food Web Interactions**

191

192 Grazing rate depends on the phytoplankton carbon concentration, which most closely represents
193 the food concentration available to the grazer (Elser and Urabe 1999; Loladze et al. 2000).

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

$$195 \quad \frac{dZ_s}{dt} = \lambda G_s - (R + G_Z + m_{1s} Z_s + m_{2s} Z_s^2) \quad (11a)$$

$$196 \quad \frac{dZ_L}{dt} = \lambda G_L - (R + m_{1L} Z_L + m_{2L} Z_L^2) \quad (11b)$$

197 where



$$198 \quad G_S = G_{S0}(1 - e^{-a_s C_{ps}})Z_S \quad (12a)$$

$$199 \quad G_L = G_{L0}(1 - e^{-a_l(C_{pl}+Z_s)})Z_L \quad (12b)$$

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

$$204 \quad G_P = G_L \frac{P_l}{P_l+Z_s} \quad (13a)$$

$$205 \quad G_Z = G_{Ll} \frac{Z_s}{P_l+Z_s} \quad (13b)$$

206 Zooplankton biomass loss to respiration is given by

$$207 \quad R = \max\{r_z T_f Z - C_{XS}, 0\} \quad (14)$$

208 and uses the same activation energy as photosynthesis. Respiration (R) is assumed to consume
 209 only carbon and not result in catabolism of existing biomass when “excess” carbon is available
 210 in the prey. In addition, conservation of mass must be maintained by recycling to the dissolved
 211 pool grazer consumption of elements in excess of biosynthetic requirements when grazer and
 212 prey elemental ratios differ. In the case where the nutrient quota (relative to carbon) exceeds the
 213 grazer fixed ratio, the excess nutrient is remineralized to the dissolved inorganic pool. In the case
 214 where the nutrient quota is less than the grazer ratio, the grazer intake is reduced to what can be
 215 supported by the least abundant nutrient (relative to the grazer biomass ratio) and excess carbon
 216 is remineralized. For the case of two nutrients (in this case N and Fe) it is necessary to define

$$217 \quad G' = G \min \left\{ \frac{N_p}{C_p} R_{CN}, \frac{Fe_p}{C_p} R_{CFe}, 1 \right\} \quad (15)$$



218 where G is equal to G_S (equation 12a) for small zooplankton and G_P (equation 13a) for large
 219 zooplankton, and R_{XY} indicates the fixed ratio of element X to element Y in grazer biomass. The
 220 'excess' carbon available for respiration is

$$221 \quad C_{XS} = G' \left\{ \frac{C_P}{N_P} R_{NC} - 1, \frac{C_P}{Fe_P} R_{FeC} - 1, 0 \right\} \quad (16a)$$

222 and the excess nutrients remineralized to their inorganic pools are

$$223 \quad N_{XS} = G' \max \left\{ \frac{N_P}{C_P} - R_{NC}, 0 \right\} \varepsilon + G' \max \left\{ R_{NC} \left(\frac{N_P}{Fe_P} R_{FeN} - 1 \right), 0 \right\} (1 - \varepsilon) \quad (16b)$$

$$224 \quad Fe_{XS} = G' \max \left\{ \frac{Fe_P}{C_P} - R_{FeC}, 0 \right\} \varepsilon + G' \max \left\{ R_{FeC} \left(\frac{Fe_P}{N_P} R_{NFe} - 1 \right), 0 \right\} (1 - \varepsilon) \quad (16c)$$

225 where

$$226 \quad \varepsilon = \frac{\max\{C_{XS}, 0\}}{C_{XS} + \Delta}$$

227 is a switch to prevent double-counting in cases where one of the terms is redundant (the excess
 228 relative to the least abundant element is included in the other term), but would otherwise be
 229 nonzero (Δ is a constant equal to 10^{-15} , to prevent divide-by-zero). For three elements, there are
 230 $3! = 6$ possible cases: for N greater or less than $C_P R_{NC}$, Fe may be either in excess relative to
 231 both C and N, deficient relative to both, or in excess relative to one but not the other (Table 2).

232

233 Table 2 - Cases where the 'excess' terms are nonzero. These terms are always greater than or
 234 equal to zero, and always zero when the phytoplankton elemental ratio is equal to the grazer bio-
 235 mass ratio. A plus (+) sign indicates that a specific term is positive. N_1 and N_2 , Fe_1 and Fe_2 indi-
 236 cate the first and second terms in equations 16b and 16c. R_{NC} is the grazer N/C (Redfield) ratio.
 237

	Fe in excess relative to both C and N					Fe in excess relative to C or N but not both					Fe deficient relative to both C and N				
	C	N_1	N_2	Fe_1	Fe_2	C	N_1	N_2	Fe_1	Fe_2	C	N_1	N_2	Fe_1	Fe_2
$N/C > R_{NC}$		+		+			+		+		+		+		
$N/C < R_{NC}$	+				+	+				+	+		+		



238 2.3 Organic and Inorganic Pools

239

240 There are two pools of detritus with different sinking rates but the same fixed elemental ratios.

241 Detrital C/N/Fe ratios are the same as zooplankton, so zooplankton mortality or grazing of small

242 zooplankton by large zooplankton produce no 'excess'. Phytoplankton mortality, and defecation

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

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

245 of grazing on phytoplankton.

246 The conservation equations for detrital C are

$$247 \frac{dD_s}{dt} = m_1(C_{ps} + Z_s) + m_2(C_{ps}^2 + Z_s^2) - r_1 D_s T_g - w_s \frac{dD_s}{dz} \quad (17a)$$

$$248 \frac{dD_l}{dt} = m_1(C_{pl} + Z_l) + m_2(C_{pl}^2 + Z_l^2) - r_2 D_l T_g - w_l \frac{dD_l}{dz} \quad (17b)$$

249 where T_g is an Arrhenius function for temperature dependence of remineralization and w is the

250 sinking speed. The conservation equations for inorganic C, N, and Fe are

$$251 \frac{dC_i}{dt} = (\zeta V^N - P_{\text{phot}}^C) C_p + R + C_{XS} + (r_1 D_s + r_2 D_l) T_g \quad (18a)$$

$$252 \frac{dN_i}{dt} = -\frac{V^N}{Q^N} N_p \left(\frac{L_{\text{NO}_3}}{L_{\text{NO}_3} + L_{\text{NH}_4}} \right) + N_{\text{ox}} - N_{\text{dentr}} (1 - A_f) \quad (18b)$$

$$253 \frac{dN_a}{dt} = -\frac{V^N}{Q^N} N_p \left(\frac{L_{\text{NH}_4}}{L_{\text{NO}_3} + L_{\text{NH}_4}} \right) + \frac{R}{R_{\text{CN}}} + N_{\text{XS}} + (r_1 D_s + r_2 D_l) R_{\text{NC}} T_g - N_{\text{ox}} + N_{\text{dnf}} - N_{\text{dentr}} A_f \quad (18c)$$

$$254 \frac{dFe}{dt} = \frac{V^{\text{Fe}}}{Q^{\text{Fe}}} Fe_p + \frac{R}{R_{\text{CFe}}} + Fe_{\text{XS}} + (r_1 D_s + r_2 D_l) R_{\text{FeC}} T_g \quad (18d)$$

255 where N_{ox} is microbial oxidation of ammonium to nitrate (nitrification), N_{dnf} and N_{dentr} are

256 sources and sinks associated with dinitrogen fixation and denitrification, and A_f is the ammonium

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



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

$$260 \quad \frac{dO_2}{dt} = -\frac{dC_i}{dt} + 2\frac{V^N}{Q^N}N_p\left(\frac{L_{NO_3}}{L_{NO_3}+L_{NH_4}}\right) - 2N_{ox} \quad (19)$$

261 Nitrification is given by

$$262 \quad N_{ox} = k_{NH_4ox}N_a\frac{K_E}{K_E+E(z)} \quad (20)$$

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

$$266 \quad N_{dnf} = k_{dnf}T_{dnf}(1 - e^{-aE})\left(\frac{Fe}{K_{Fe}+Fe}\right)\left(\frac{K_{NO_3}}{K_{NO_3}+N_i+N_a}\right) \quad (21)$$

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

269

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

$$272 \quad N_{frxn} = 1 - \frac{\min(O_2, O_{mxd})}{O_{mxd}} \quad (22)$$

273 Remineralization is then divided among oxygen ($1 - N_{frxn}$), nitrate ($0.875N_{frxn}$), and ammonium
274 ($0.125N_{frxn}$) assuming an average anammox contribution of 25% (Babbin et al., 2014). We use
275 this average ratio of anammox to classical denitrification to partition fixed N losses between
276 NO_3^- and NH_4^+ ; the DIC sink and organic matter source associated with anammox are small and
277 are neglected here.

278



279 2.4 Calcification, Calcite Dissolution, and Alkalinity

280

281 Calcification is represented by a detrital calcium carbonate (CaCO_3) state variable, but no
282 explicit calcifier groups. Detrital CaCO_3 sinks in the same fashion as detrital particulate organic
283 carbon (POC), with a sinking rate independent of those for large and small organic detritus.

284 Calcite production is represented as a fixed fraction of detritus production from small
285 phytoplankton and small zooplankton mortality:

$$286 \quad \frac{dCa}{dt} = m_1(C_{ps} + Z_s)P_{Ca} + m_2(C_{ps}^2 + Z_s^2)P_{Ca} - k_{Ca}Ca - w_{Ca}\frac{dCa}{dz} \quad (23)$$

287 Calcite dissolution occurs throughout the water column as a first order process (i.e., no
288 dependence on temperature or saturation state). Approximately 80% of calcite produced is
289 exported from the euphotic zone. Burial in the sediments is represented as a simple 'on/off'
290 switch dependent on the calcite saturation state (zero when $\Omega_C < 1$ and 1 when $\Omega_C \geq 1$). Calcite
291 burial is balanced by an equivalent source of DIC and alkalinity at the ocean surface as a crude
292 parameterization of fluvial sources.

293

294 For each mole of calcite production two moles of alkalinity equivalent are lost from the
295 dissolved phase; the reverse occurs during calcite dissolution. There are additional sources and
296 sinks for alkalinity associated with phytoplankton nutrient uptake, organic matter
297 remineralization, nitrification, denitrification and dinitrogen fixation (Wolf-Gladrow et al., 2007,
298 see Supplementary Table S2). The anammox reaction does not in itself contribute to alkalinity
299 (Jetten et al., 2001), but there is a sink associated with ammonium oxidation to nitrite (the model
300 does not distinguish between nitrite and nitrate). Autotrophic production of organic matter by
301 anammox bacteria is a net source of alkalinity (Strous et al., 1998) but this source is extremely



302 small (~ 0.03 mol/molN) and is neglected here. Globally, the sources and sinks of alkalinity from
303 the N cycle offset each other such that there is no net gain or loss as long as the global fixed N
304 pool is conserved (see below Sect. 2.5). If dinitrogen fixation and denitrification are allowed to
305 vary freely, there will generally be a net gain or loss of fixed N and, therefore, of alkalinity.

306

307 **2.5 External Nutrient Sources and Sinks**

308

309 External sources and sinks consist of river inputs, aeolian deposition, biological N₂ fixation, de-
310 nitrification, mobilization of Fe from reducing sediments, loss of Fe to scavenging, and burial of
311 calcium carbonate in the sediments. Aeolian deposition of Fe is calculated from a climatology of
312 mineral dust deposition generated from offline (atmosphere-only) simulations with CanAM4
313 (von Salzen et al., 2013), with an Fe mass fraction of 5% and a fractional solubility of 1.4% in
314 the surface layer. Subsurface dissolution is parameterized based on PISCES (Aumont et al.,
315 2015); the total dissolution is 6.35%, with 22% of soluble Fe input into the first vertical layer
316 (see Supplementary material). Iron from reducing sediments is also based on PISCES, with a
317 constant areal flux of $1000 \text{ nmol m}^{-2} \text{ d}^{-1}$ in the first model level, declining exponentially with in-
318 creasing seafloor depth (i.e., assuming that the sediments become progressively more oxygen-
319 ated) with an e-folding length scale of about 200 m. Scavenging of dissolved iron is first-order
320 with a high rate (2.5 d^{-1}) for concentrations in excess of 0.6 nM (Johnson et al., 1997). For con-
321 centrations below this threshold, the rate is much lower (0.001 d^{-1}) and is weighted by the con-
322 centration of organic detritus (Christian et al., 2002b), i.e.,

$$323 \quad \frac{dFe}{dt} = -FeS_{Fe1} \min\{(D_S + D_L)P_{Fe}, 1\} \quad (24)$$



324 where Fe is the dissolved iron concentration, D_S and D_L are the small and large detritus concen-
325 trations, S_{Fe1} is the first-order scavenging rate in surface waters with abundant particulates, and
326 P_{Fe} is an empirical parameter to determine the dependence on particle concentration (Table 1).
327 The basis for this parameterization is that the rate of scavenging must depend not only on the
328 concentration of iron but on the concentration of particles available for it to precipitate onto, and
329 assumes that POC is strongly positively correlated with total particulate matter. Scavenging is
330 treated as irreversible, i.e., scavenged Fe is not tracked and does not reenter the dissolved phase.

331

332 Unlike in CMOC, N_2 fixation and denitrification vary independently in CanOE, so the global
333 total N pool can change. Conservation is imposed by adjusting the global total N pool according
334 to the difference between the gain from N_2 fixation and the loss to denitrification. A slight
335 adjustment is applied to the nitrate concentration at every grid point, while preserving the overall
336 spatial structure of the nitrate field. Adjustments are multiplicative rather than additive to avoid
337 producing negative concentrations. This adjustment does not maintain (to machine precision) a
338 constant global N inventory but is intended to minimize long term drift, keeping it much smaller
339 than the free surface error (see below). This adjustment is applied every 10 days and has a
340 magnitude of approximately 7×10^{-8} of the total N.

341

342 One mole of alkalinity is removed per mole of N added or removed, since there are alkalinity
343 sources of 1 mol/molN associated with both N_2 fixation (creation of new NH_4^+) and
344 denitrification (removal of NO_3^-), offset by a 2 mol/molN sink associated with nitrification. As
345 noted above, $CaCO_3$ can dissolve or be buried in the sediments depending on the calcite
346 saturation. DIC and alkalinity lost to burial are reintroduced at the ocean surface, at the same grid



347 point as burial occurs, providing a crude parameterization of river inputs so that global
348 conservation is maintained (fresh water runoff contains no DIC or alkalinity). However, the OPA
349 free surface formulation is inherently imperfect with regard to tracer conservation. Drift in total
350 ocean alkalinity and nitrogen over time is on the order of 0.01% and 0.03% per thousand years,
351 respectively (losses due to the free surface are generally larger for tracers with less homogeneous
352 distributions).

353

354 **2.6 Ancillary data**

355

356 For first-order model validation we have relied largely on global gridded data products rather
357 than individual profile data. Global gridded data from World Ocean Atlas 2013 (WOA2013)
358 (Locarnini et al., 2013; Zweng et al., 2013; Garcia et al., 2014a; 2014b) were used for
359 temperature, salinity, and oxygen and nitrate concentration. DIC and alkalinity were taken from
360 the GLODAP gridded data product (Lauvset et al., 2016). Offline carbon chemistry calculations
361 were done following the Best Practices Guide (Dickson et al., 2007) and the OMIP-BGC
362 protocols (Orr et al., 2017), which are identical to those used in the models except that constant
363 reference concentrations were used for phosphate (1 μM) and silicate (10 μM).

364

365 There is no global gridded data product for Fe, but we have made use of the GEOTRACES
366 Intermediate Data Product 2017 (Schlitzer et al., 2018), and the data compilations from MBARI
367 (Johnson et al., 1997; 2003) and PICES Working Group 22 (Takeda et al., 2013). The latter two
368 are concentrated in the Pacific, while GEOTRACES is more global. The combined data sets
369 provide more than 10000 bottle samples from more than 1000 different locations (Supplementary



370 Figure S4a) (excluding some surface transect data that involve frequent sampling of closely
371 spaced locations along the ship track). More detail about model comparison to these data
372 compilations and the list of original references are given in the Supplementary information.
373
374 Satellite ocean colour estimates of surface chlorophyll were taken from the combined
375 SeaWiFS/MODIS climatology described by Tesdal et al. (2016). Climatological satellite POC
376 was downloaded from the NASA ocean colour web site and is based on the algorithm of
377 Stramski et al. (2008) using MODIS-Aqua data. This climatology differs slightly from the
378 chlorophyll one in terms of years included and sensors utilized, but as only climatological
379 concentrations are considered and each climatology covers ~15 years, these differences will have
380 negligible effect on the results presented.
381
382 CMIP6 model data were regridded to a common grid ($2 \times 2^\circ$, 33 levels following the GLODAP
383 levels) to facilitate ensemble averaging. The years 1986-2005 of the Historical experiment were
384 averaged into climatologies or annual means, for meaningful comparison with observation-based
385 data products. A single realization was used in each case; as 20 year averages are used, internal
386 variability is assumed to have little effect (e.g., Arguez and Vose, 2011). Sampling among
387 CMIP6 models was somewhat opportunistic and the exact suite of models varies among the
388 analyses presented. When we conducted a search for a particular data field, we included in the
389 search parameters all models that published that field, and repeated the search at least once for
390 models that were unavailable the first time the search was executed. In some cases, model
391 ensemble means excluded all but one model from a particular ‘family’ (e.g., there are three
392 different MPI-ESM models for which ocean biogeochemistry fields were published), as the



393 solutions were found to be similar and would bias the ensemble mean towards their particular
394 climate. More detail is given in Supplementary Table S3.

395

396 **3. Results**

397

398 **3.1 Distribution of oxygen**

399

400 The spatial distribution of oxygen concentration ($[O_2]$) at selected intermediate depths (400, 900,
401 and 1400 m) for CanESM5, CanESM5-CanOE, a model ensemble mean (MEM) of CMIP6
402 models (excluding CanESM5 and CanESM5-CanOE), and gridded data from WOA2013 is
403 shown in Figure 2. The major features are consistent across the models. All three cases show
404 elevated oxygen concentrations relative to observations, particularly in the North Pacific, the
405 North Atlantic and the Southern Ocean. In the Indian Ocean, both CanESM models show high
406 oxygen concentrations in the Arabian Sea and deeper layers of the Bay of Bengal relative to
407 observations and other CMIP6 models; these biases are somewhat smaller in CanESM5-CanOE
408 than in CanESM5 (Figure 2).

409

410 Biases in the eastern boundary current regions are depth and model specific. CanESM5 shows
411 particularly strong oxygen depletion at 1400 m in the eastern tropical Pacific. In the southeastern
412 Atlantic, models tend to be biased low at the shallower depths, and show somewhat more
413 variation at greater depths (Figure 2). Overall, $[O_2]$ biases tend to be positive over large areas of
414 ocean with the exception of some eastern boundary current regions, implying that models
415 exaggerate the extent to which remineralization is concentrated in these regions. An alternate

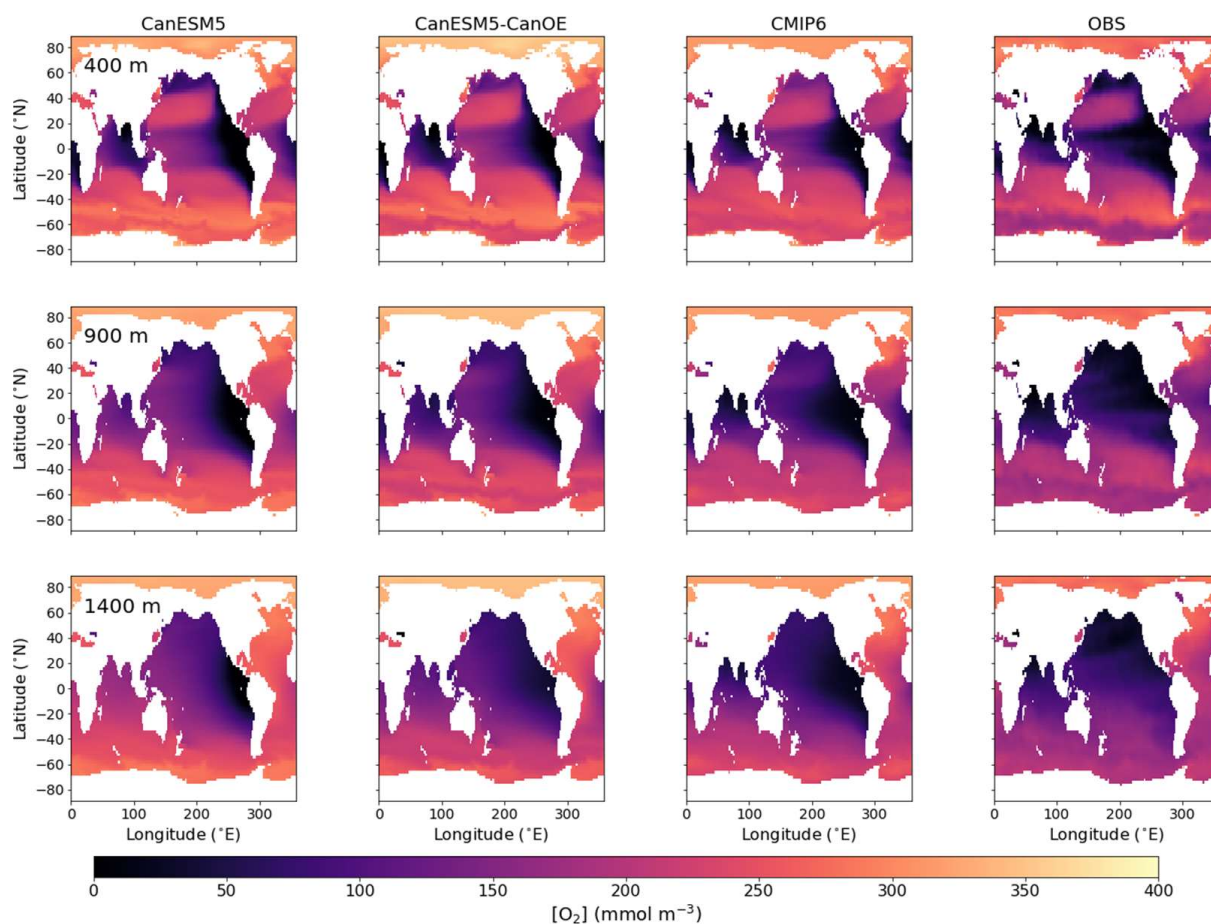


Figure 2 - Global distribution of oxygen (O_2) concentration in mmol m^{-3} at 400, 900, and 1400 m (rows) for CanESM5-CanOE, CanESM5, the mean for other (non-CanESM) CMIP6 models, and World Ocean Atlas 2013 (WOA2013) observations (columns). Difference from the observation-based fields are shown in Supplementary Figure S3.



417 version of Figure 2 that shows model errors relative to the observational data product is given in
418 Supplementary Figure S3.

419

420 The zonal mean oxygen concentration, saturation concentration, and apparent oxygen utilization
421 (AOU) are shown in Figure 3 for the same four cases. Again, the models generally show a
422 positive bias in $[O_2]$, particularly in high-latitude deep waters. The major ocean circulation
423 features are reproduced fairly well in all cases (e.g., weaker ventilation of low-latitude
424 subsurface waters, greater vertical extent of well-ventilated surface waters in the subtropics). The
425 saturation concentration (a function of temperature and salinity) generally shows relatively little
426 bias, implying that the bias in $[O_2]$ arises mainly from remineralization and/or ventilation. AOU
427 is lower than observed over much of the subsurface ocean. Regional biases are quite consistent
428 across models, but are slightly greater in CanESM5 than in CanESM5-CanOE or the ensemble
429 mean, except in the Arctic Ocean. Again, Supplementary Figure S3 includes a version of this
430 plot that shows the model differences from the observations.

431

432 The skill of each model with respect to the distribution of O_2 at different depths is represented by
433 Taylor diagrams (Taylor, 2001) in Figure 4, in which all of the CMIP6 models that were shown
434 as an ensemble mean in Figures 2 and 3 are shown individually. The blue dots represent
435 CanESM5, red CanESM5-CanOE, and grey the ensemble mean of all CMIP6 models except
436 CanESM5 and CanESM5-CanOE; the smaller grey dots represent the individual models.
437 CanESM5-CanOE shows slightly higher pattern correlation than CanESM5 at all depths. Both
438 models compare favourably with the full suite of CMIP6 models, with $r > 0.85$ for CanESM5 and
439 $r > 0.9$ for CanESM5-CanOE at all depths examined, and a normalized standard deviation within

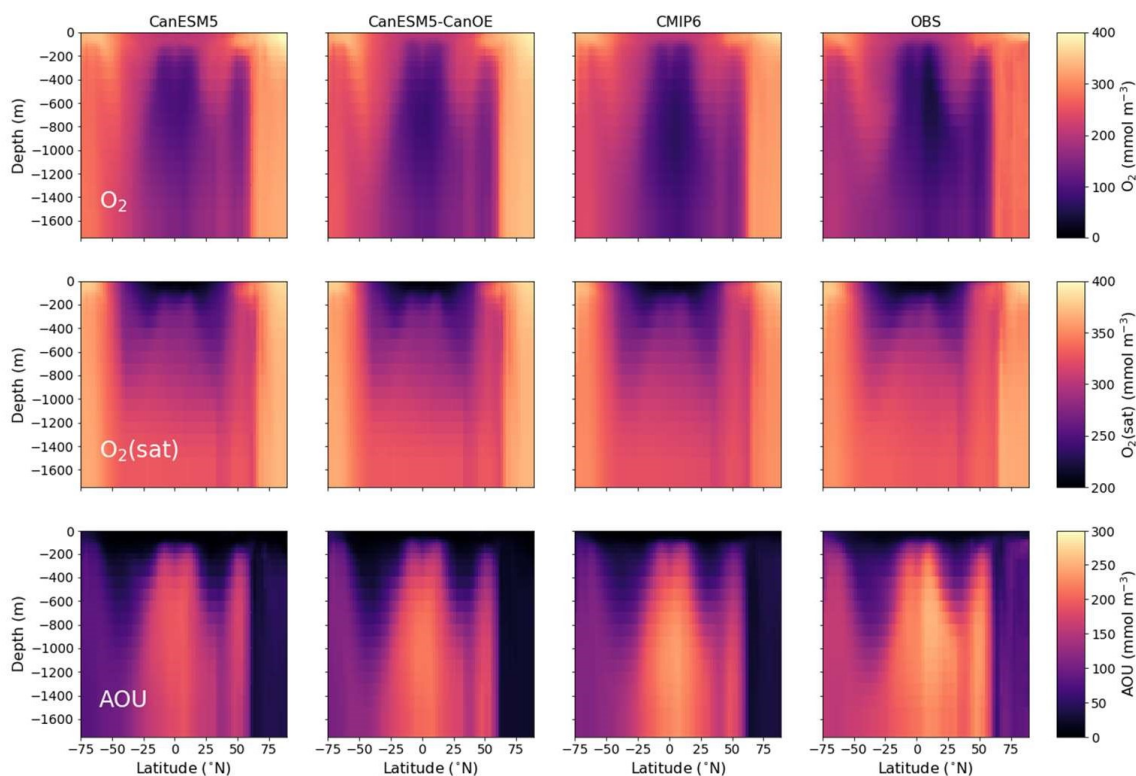


Figure 3 - Latitude-depth distribution (surface to 1750 m) of zonal mean oxygen concentration (O_2), oxygen concentration at saturation ($O_2(\text{sat})$), and apparent oxygen utilization (AOU) in mmol m^{-3} for CanESM5-CanOE, CanESM5, the mean for other CMIP6 models, and observations (WOA2013). Note different colour scales for different rows. Difference from the observation-based fields are shown in Supplementary Figure S3.

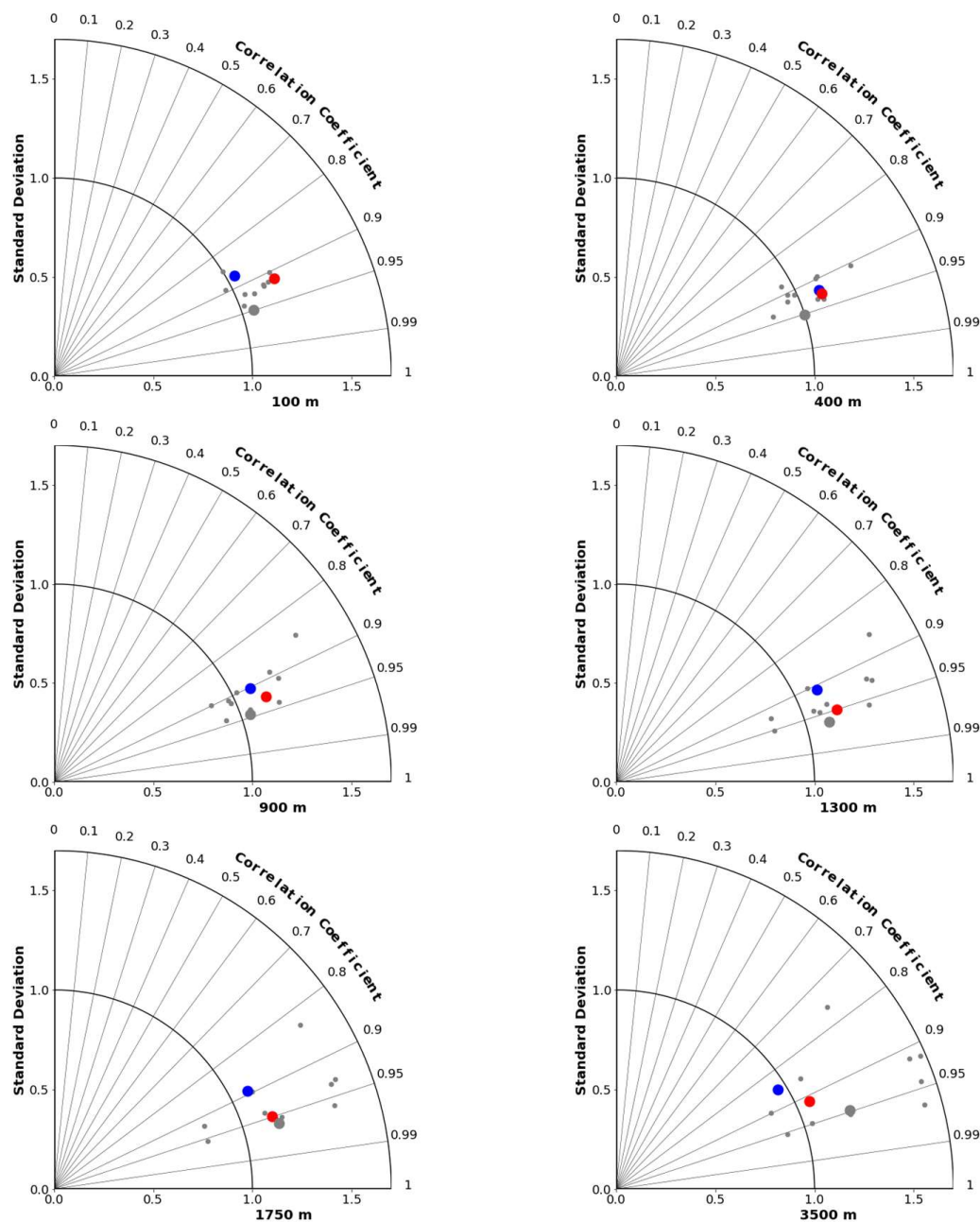


Figure 4 - Taylor diagrams (Taylor, 2001) comparing modelled and observed distributions of oxygen at specific depths from 100 to 3500 m. Angle from the vertical indicates spatial pattern correlation. Distance from the origin indicates ratio of standard deviation in modelled vs. observed (WOA2013) fields. 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.



442 $\pm 25\%$ of unity.

443

444 The total volume of ocean with $[O_2]$ less than 6 mmol m^{-3} (the threshold for denitrification
445 (Devol, 2008)) and 60 mmol m^{-3} (a commonly used index of hypoxia) is shown in Figure 5. The
446 total volume is highly variable among models (note, however, that there are several clusters of
447 related models with quite similar totals). CanESM5 and CanESM5-CanOE have among the
448 lowest total volumes (i.e., the interior ocean is relatively well ventilated) and are among the
449 nearest to the observed total. For $[O_2] < 60 \text{ mmol m}^{-3}$ the bias is, nonetheless, quite large (i.e., the
450 observed volume is underestimated by almost 50% in both models). The volume of water with
451 $[O_2]$ below the denitrification threshold is overestimated in both CanESM5 and CanESM5-
452 CanOE; CanESM5-CanOE has a much smaller total that is closer to the observed value. The bias
453 in the spatial pattern of hypoxia (not shown) is generally similar to the bias in dissolved oxygen
454 distribution (Figure 2). The low-oxygen regions are generally more concentrated in the eastern
455 tropical Pacific in the models than in observations, and the low-oxygen region in the northwest
456 Pacific is not well reproduced in CanESM models.

457

458 **3.2 Distribution of DIC, alkalinity, and $CaCO_3$ saturation**

459

460 The spatial distribution of aragonite saturation state (Ω_A) at selected depths is shown in Figure 6
461 (the first two depths are the same as in Figure 2, but the third is much deeper). In this case the
462 observations are a combination of GLODAP (Lauvset et al., 2016) for DIC and alkalinity, and
463 WOA2013 for temperature and salinity. CanESM5 and CanESM5-CanOE generally compare
464 well with other models and observations. The low saturation bias in the eastern tropical Pacific is

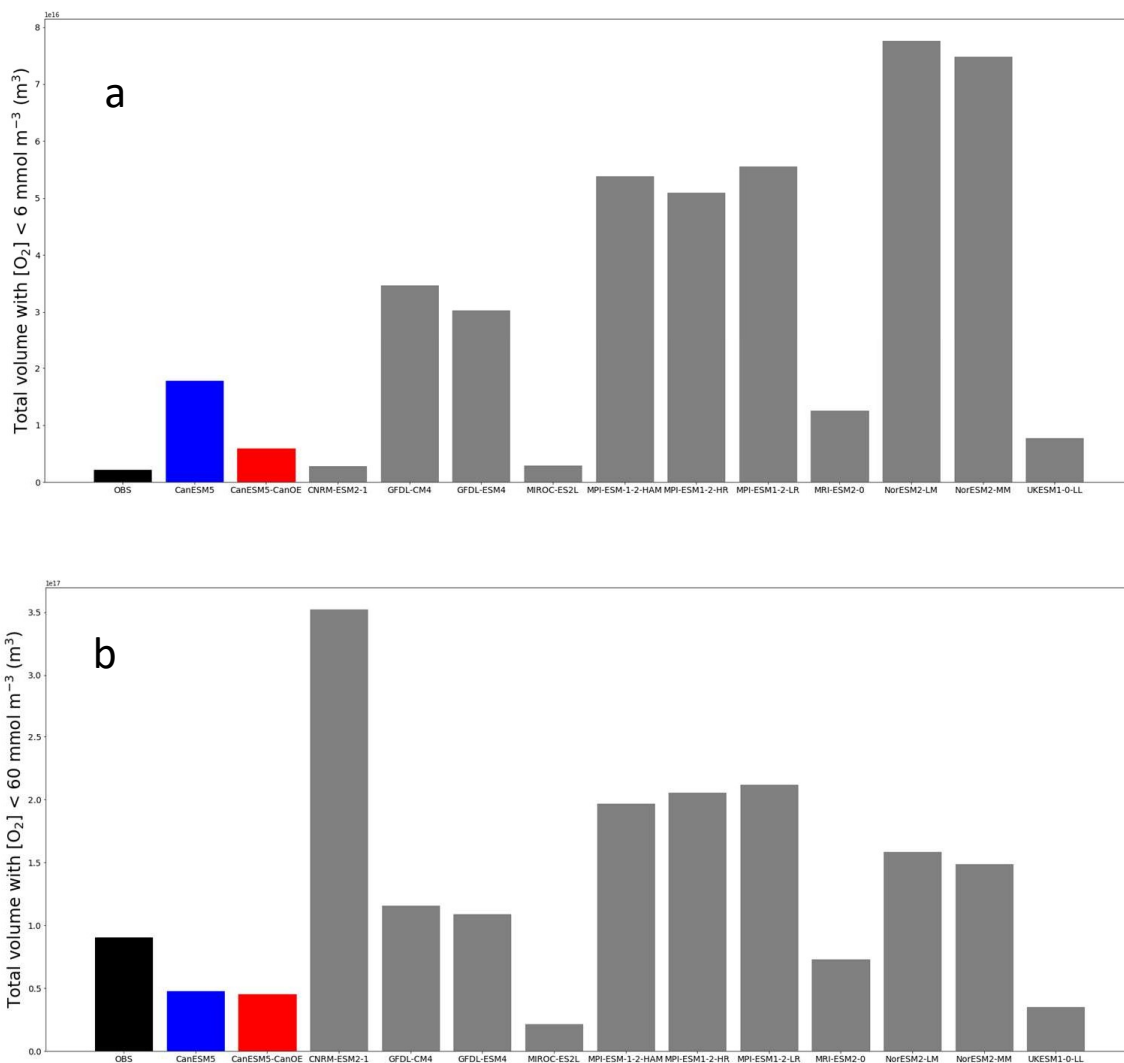


Figure 5 - Total volume of ocean with oxygen (O₂) concentration less than (a) 6 mmol m⁻³ (mean for last 30 years of the historical experiment) and (b) 60 mmol m⁻³. Observation are from WOA2013.

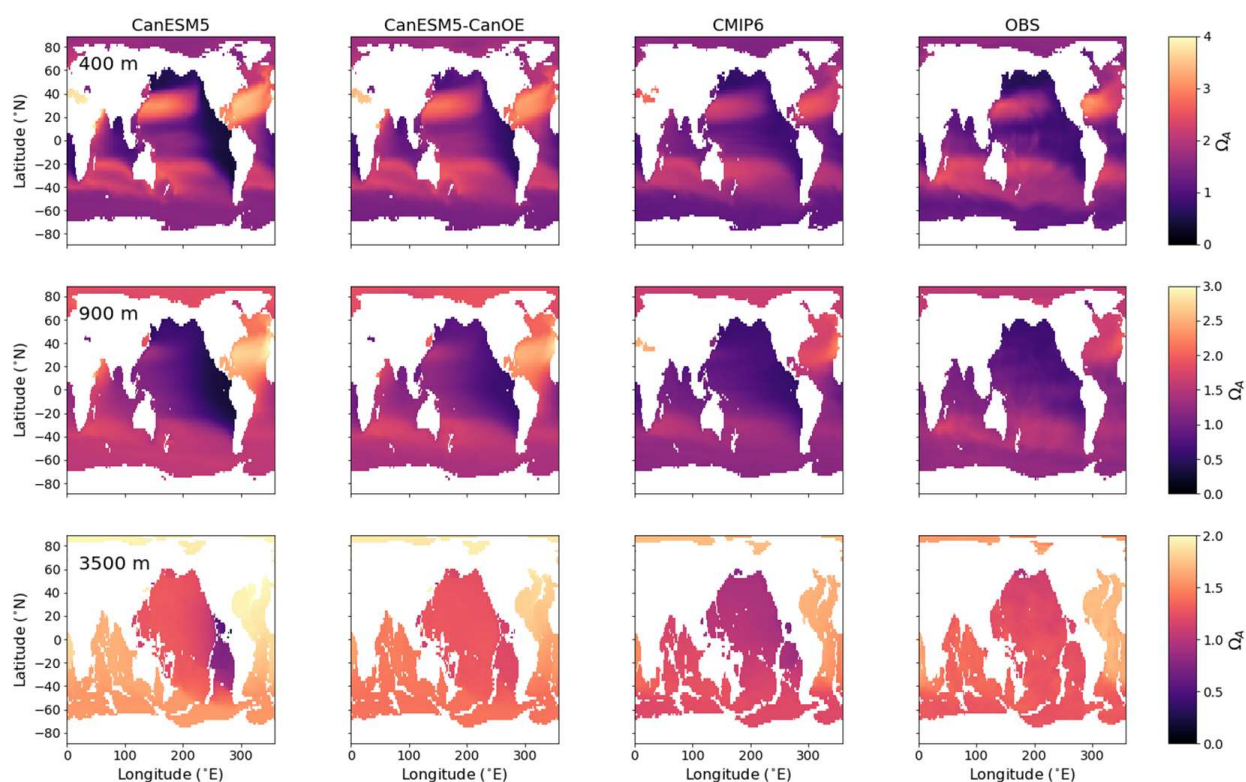


Figure 6 - Global distribution of aragonite saturation (Ω_A) at 400, 900, and 3500 m for CanESM5-CanOE, CanESM5, the mean for other CMIP6 models, and observations (GLODAP + WOA2013). Note different colour scales for different depths. Difference from the observation-based fields are shown in Supplementary Figure S3.



467 substantially reduced in CanESM5-CanOE compared to CanESM5. On the other hand CanESM5
468 generally does better than CanESM5-CanOE, or the ensemble mean, at reproducing the low
469 saturation states in the northwestern Pacific and the Bering Sea. Both models show a high
470 saturation state bias in the North Atlantic and the well-ventilated regions of the north Pacific
471 subtropical gyre. These biases are reduced slightly in CanESM5-CanOE, probably due to the
472 smaller average remineralization length scale for organic detritus.

473

474 Zonal mean distributions of aragonite saturation state (Ω_A), calcite saturation state (Ω_C), and
475 carbonate ion concentration ($[CO_3^{2-}]$) are shown in Figure 7 (Supplementary Figure S3 includes
476 versions of Figures 6 and 7 that explicitly show the model differences from the observations).
477 The models generally compare well with the observations in the representation of the
478 latitude/depth distribution of high and low saturation waters. CanESM5 has a high saturation bias
479 in low-latitude surface waters that is somewhat reduced in CanESM5-CanOE. Both CanESM5
480 models show a high saturation bias in Northern Hemisphere intermediate (e.g., 200-1000 m)
481 depth waters that is larger than in the ensemble mean.

482

483 Taylor diagrams for a range of depths are shown for DIC in Figure 8 and for Ω_A in Figure 9 (for
484 alkalinity, see Supplementary Figure S2). As expected, the MEM generally compares favourably
485 with the individual models (e.g., Lambert and Boer, 2001). CanESM5 and CanESM5-CanOE
486 compare favourably with the full suite of CMIP6 models. CanESM5-CanOE shows a gain in
487 skill relative to CanESM5, and both show improvement relative to CanESM2. At 400 m,
488 CanESM2 stands out as having extremely high variance, which is mostly due to extremely high
489 DIC concentrations occurring over a limited area in the eastern equatorial Pacific (not shown).

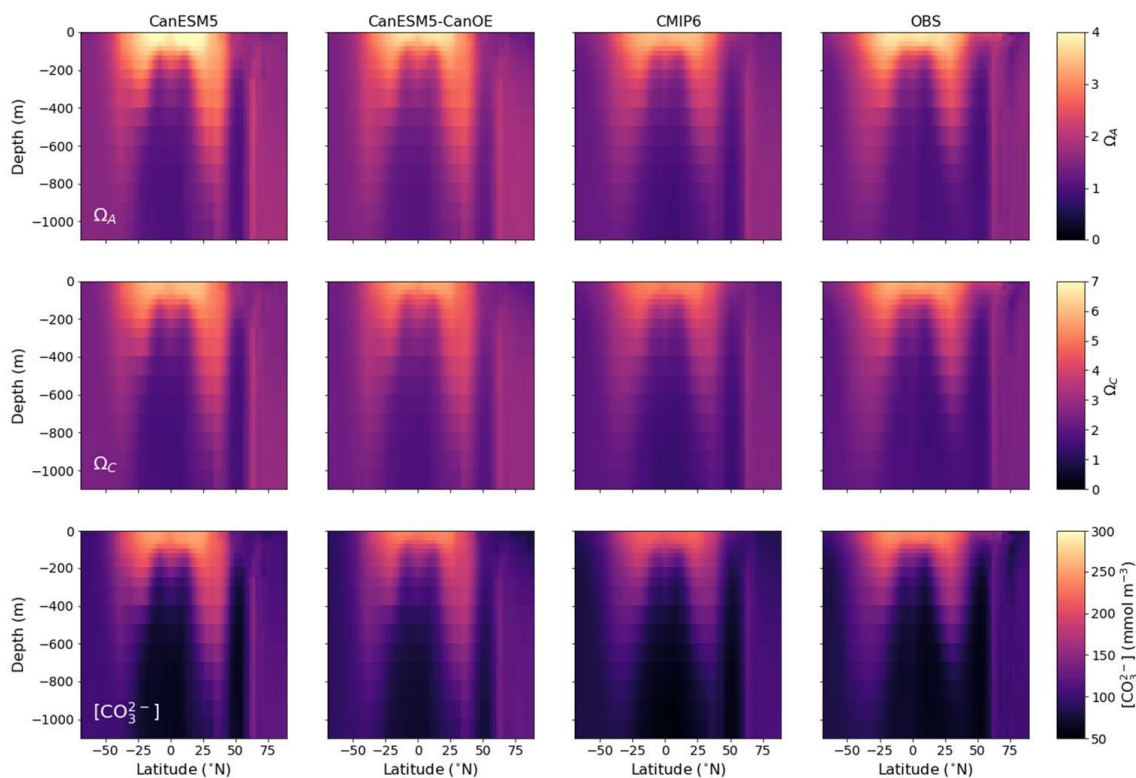


Figure 7 - Latitude-depth distribution of zonal mean (surface to 1150 m) aragonite saturation state (Ω_A), calcite saturation state (Ω_C), and carbonate ion concentration ($[\text{CO}_3^{2-}]$) in mmol m^{-3} for CanESM5-CanOE, CanESM5, the mean for other CMIP6 models, and observations (GLODAP + WOA2013). Difference from the observation-based fields are shown in Supplementary Figure S3.

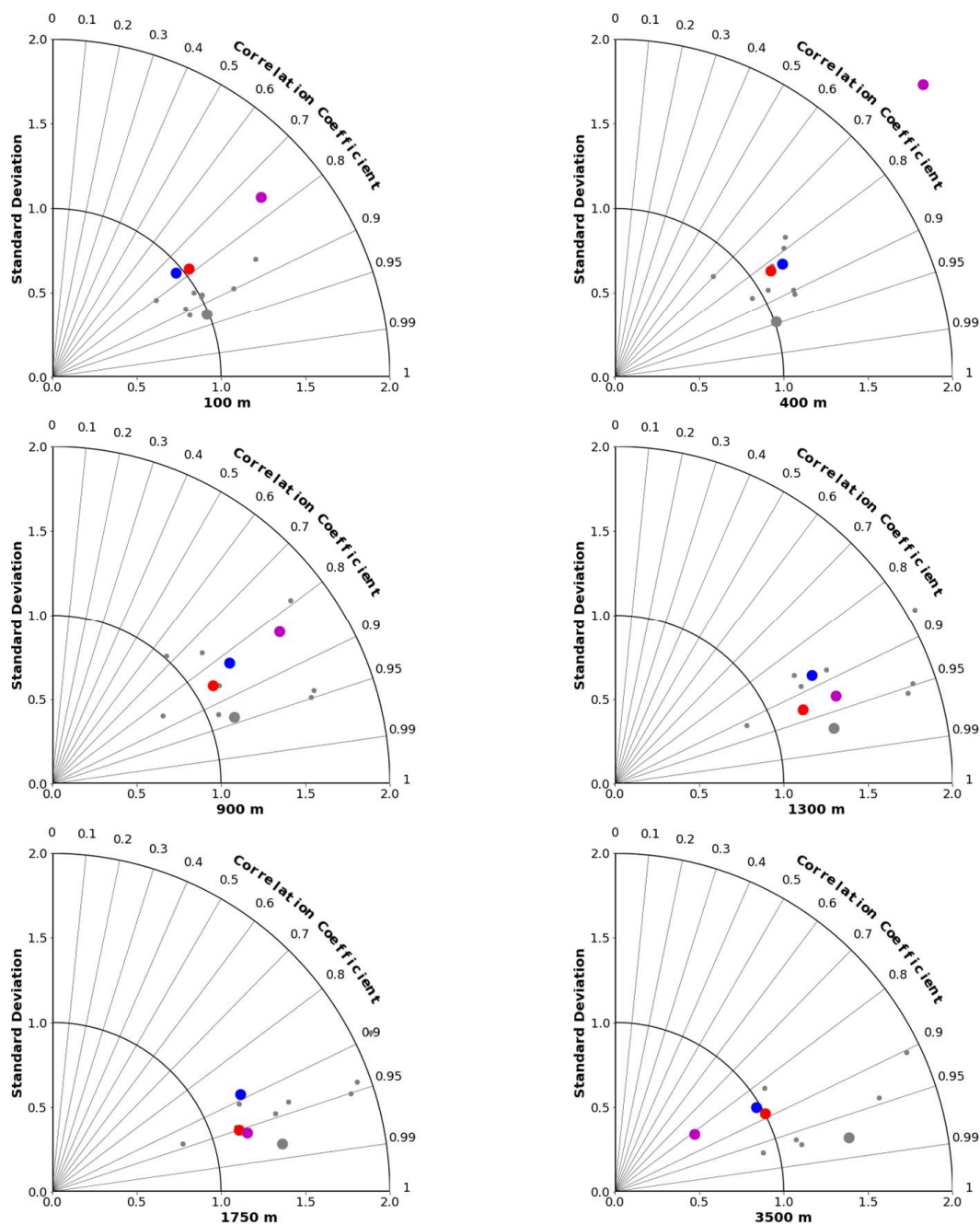


Figure 8 - Taylor diagrams comparing modelled and observed distributions of DIC at specific depths from 100 to 3500 m. Observations are from GLODAP (Lauvset et al., 2016). Red dots represent CanESM5-CanOE, blue dots CanESM5, magenta dots CanESM2, small grey dots other CMIP6 models, and large grey dots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

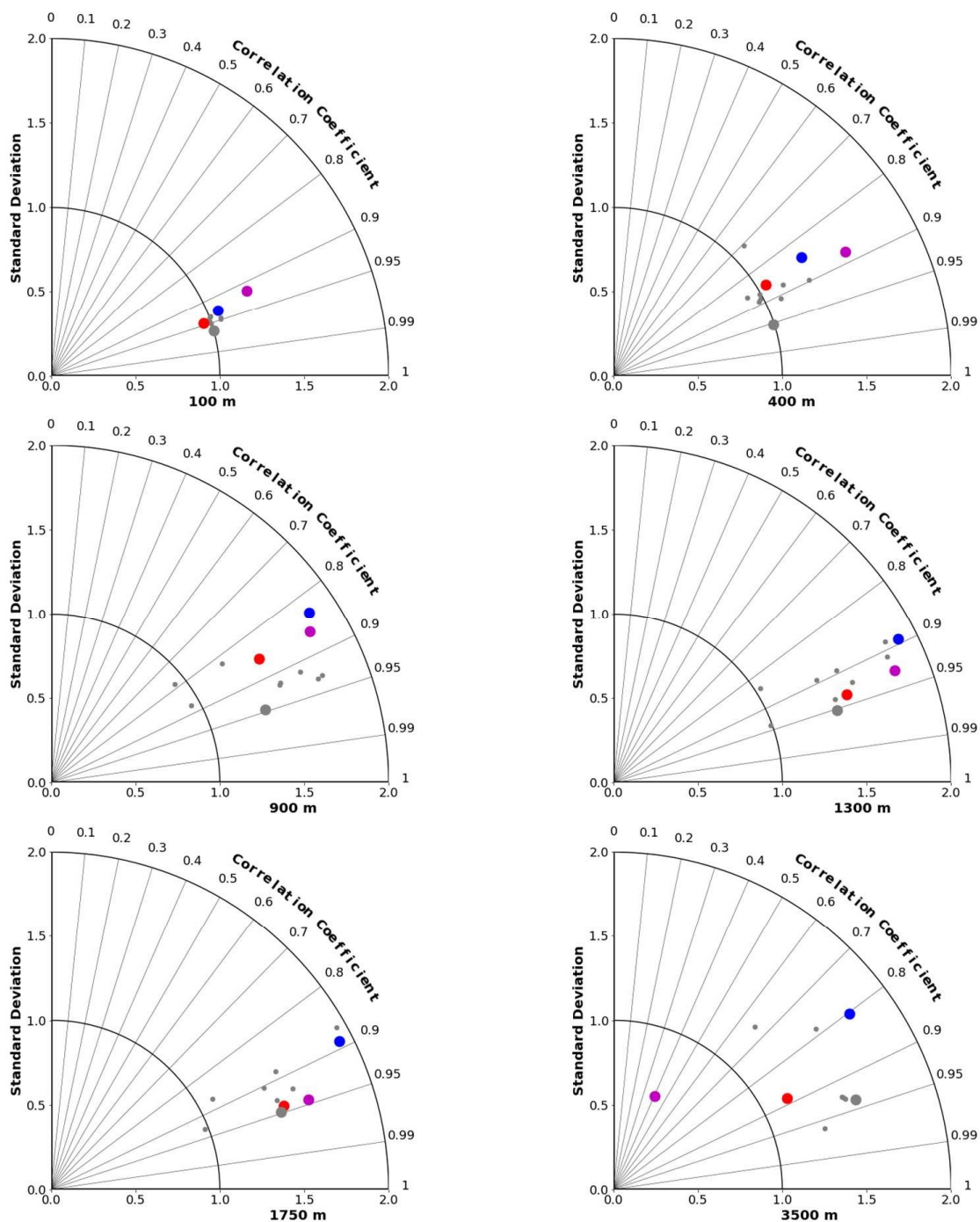


Figure 9 - Taylor diagrams comparing modelled and observed (GLODAP + WOA2013) distributions of Ω_A at specific depths from 100 to 3500 m. Symbol colours as in Figure 8.



493 This bias is present in CanESM5 and in CMIP6 models generally (Figure 6) but involves much
494 lower concentrations spread over a larger area.

495

496 **3.3 N and Fe cycles**

497

498 An important difference between CMOC and CanOE is the inclusion of a prognostic Fe cycle.
499 The CMOC iron mask (Zahariev et al., 2008) was a pragmatic solution in the face of resource
500 limitations but is inherently compromised as it can not evolve with a changing climate. Other
501 centres that introduced a prognostic Fe cycle between CMIP5 and CMIP6 include JAMSTEC
502 (MIROC-ESM, MIROC-ES2L) and the UK Met Office (HadGEM2-ES, UKESM1-0-LL). The
503 first order test of a model with prognostic, interacting Fe and N cycles is whether it can
504 reproduce the distribution of High-Nutrient, Low-Chlorophyll (HNLC) regions and the
505 approximate surface macronutrient concentrations within these. CanESM5-CanOE succeeded by
506 this standard, although the surface nitrate concentrations are biased low in the subarctic Pacific
507 and equatorial Pacific and high in the Southern Ocean and in the global mean (Figure 10).

508

509 The seasonal cycle of the zonal mean surface nitrate concentration for a selection of CMIP6
510 models is shown in Figure 11. CanESM5, CanESM5-CanOE, and CNRM-ESM2-1 reproduce the
511 equatorial enrichment and the low concentrations in the tropical-subtropical latitudes fairly well.
512 Some models either have very weak equatorial enrichment (MPI-ESM1-2-LR) or too high a
513 concentration in the off-equatorial regions (UKESM1-0-LL, NorESM2-LM). UKESM1-0-LL
514 has very high concentrations throughout the low-latitude Pacific, which biases the ensemble
515 mean (Figure 11a). Figure 11b shows the same data as Figure 11a but for a more limited latitude

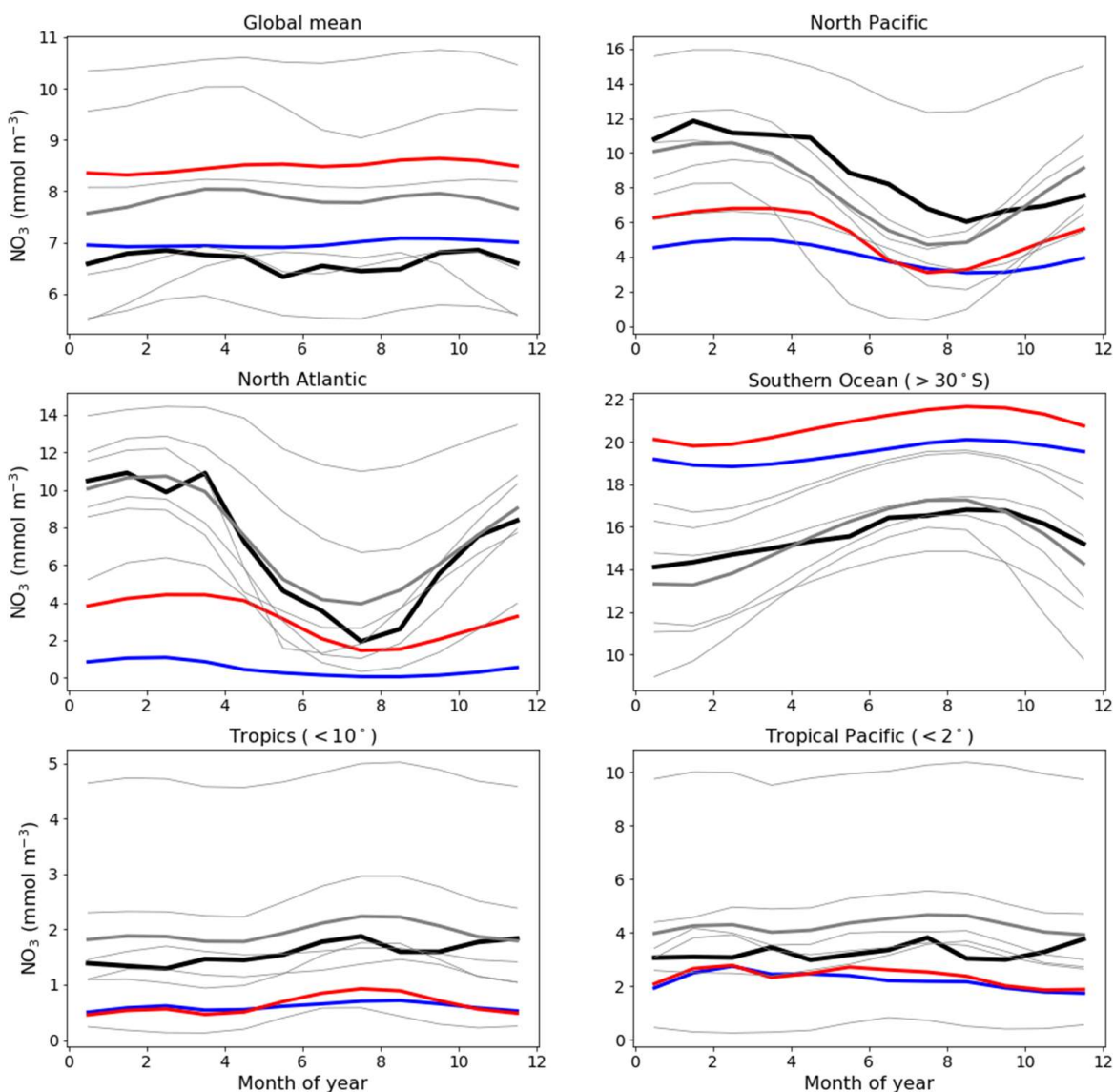


Figure 10 - Climatological seasonal cycle of surface nitrate concentration averaged for selected ocean regions. Thick red line represents CanESM5-CanOE, thick blue line CanESM5, thick black line observations (WOA2013), thin grey lines individual CMIP6 models, and thick grey line the model ensemble mean (excluding CanESM5 and CanESM5-CanOE). Regional boundaries are given in Supplementary Table S4.

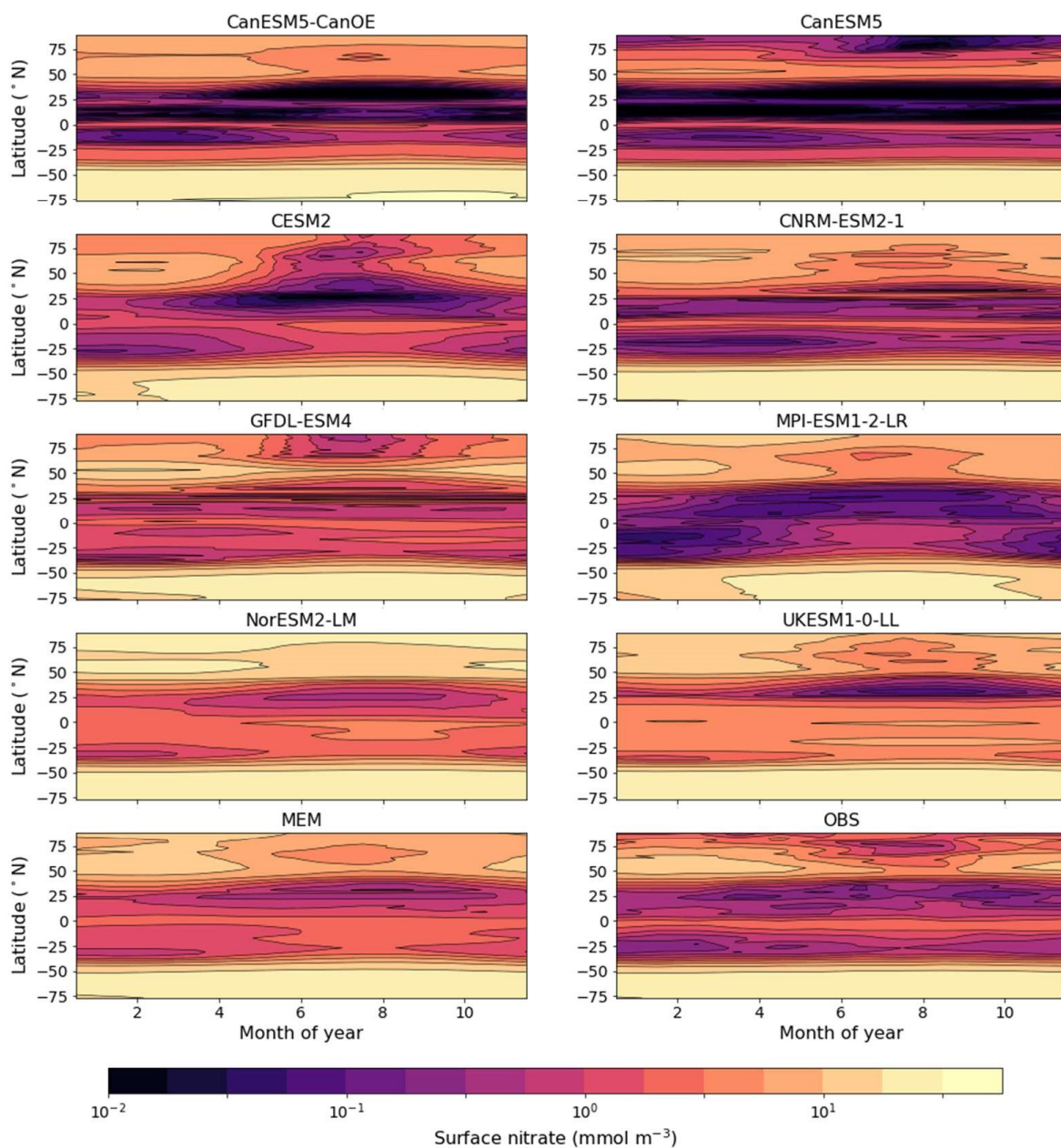


Figure 11 - (a) Climatological seasonal cycle of zonal mean surface nitrate concentration, for a selection of CMIP6 models, a model ensemble mean (MEM) excluding CanESM5 and CanESM5-CanOE, and an observation-based data product (WOA2013).

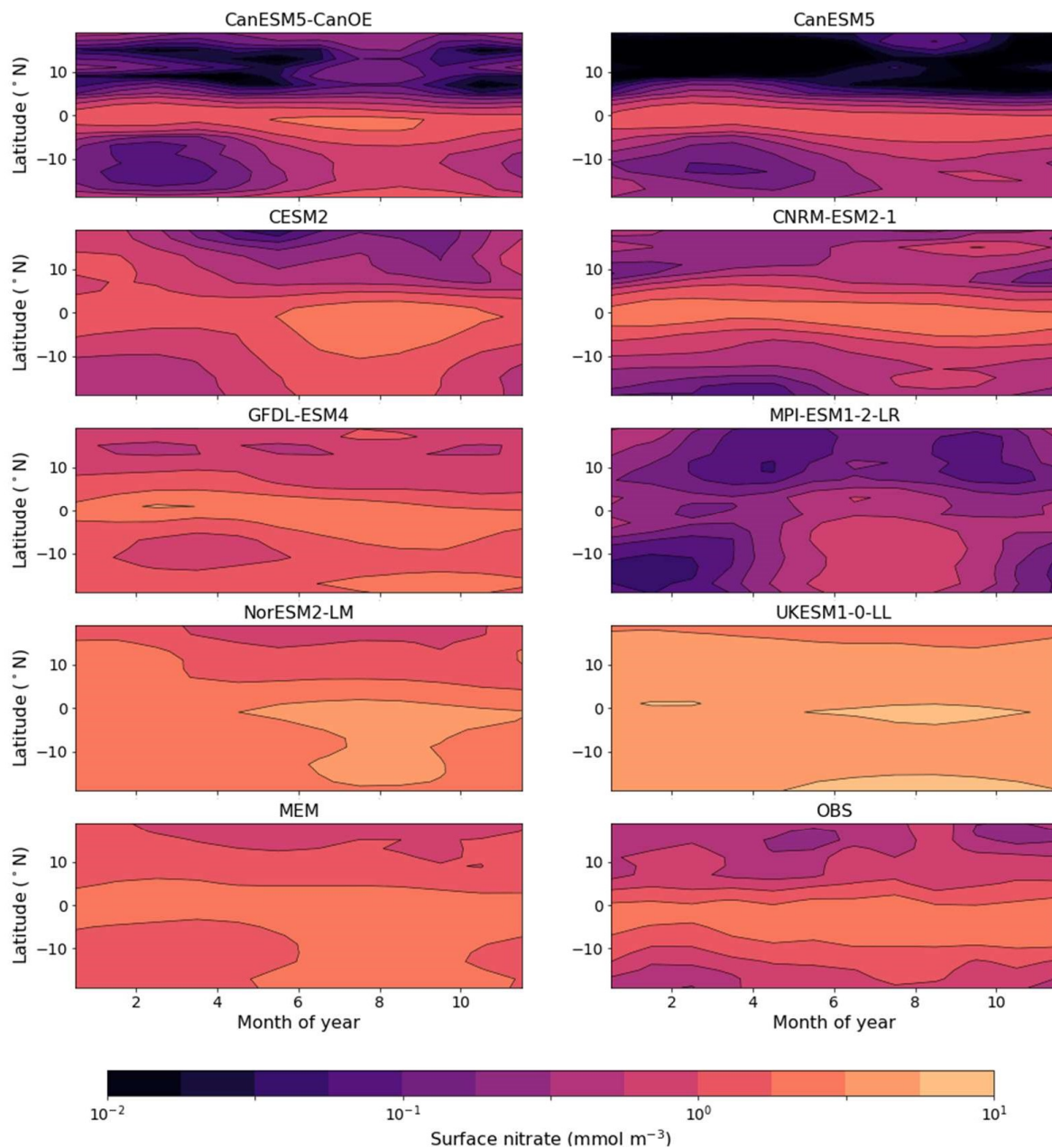


Figure 11 (b) As Figure 11(a) but for latitudes <20°.



519 range to better illustrate model behaviour in the tropics. CanESM5, CanESM5-CanOE, and
520 CNRM-ESM2-1 reproduce the seasonal cycle of tropical upwelling (e.g., Philander and Chao,
521 1991), with highest concentrations in summer.

522

523 The surface distribution of dissolved iron (dFe) in various CMIP6 models is shown in Figure 12.
524 For Fe there is no observation-based global climatology with which to compare the model
525 solutions (some comparisons to available profile data are shown in Supplementary Figures S4b-
526 g). CanESM5-CanOE shows a similar overall spatial pattern to other models, and generally falls
527 in the middle of the spread, particularly regarding concentrations in the Southern Ocean. Several
528 models show extremely high concentrations in the tropical-subtropical North Atlantic (Sahara
529 outflow region). CanESM5-CanOE, along with CNRM-ESM2-1 and CESM2, has much less
530 elevated concentrations in this region, due to lower deposition or greater scavenging or both.

531 CanESM5-CanOE has its lowest concentration in the eastern subtropical South Pacific, which is
532 common to many models (Figure 12). The area of strong surface depletion is generally more
533 spatially restricted in CanESM5-CanOE than in other models, and surface dFe concentrations are
534 greater over large areas of the Pacific. Both the north-south and east-west asymmetry of
535 distribution in the Pacific is greater in CanESM5-CanOE than in most other models, some of
536 which show the South Pacific minimum extending westward across the entire basin, and others
537 into the Northern Hemisphere. Only in CESM2 is this minimum similarly limited to the
538 southeast Pacific.

539

540 The mean depth profiles of dFe are shown in Figure 13. Some models show more of a “nutrient-
541 type” (increasing with depth due to strong near-surface biological uptake and subsequent

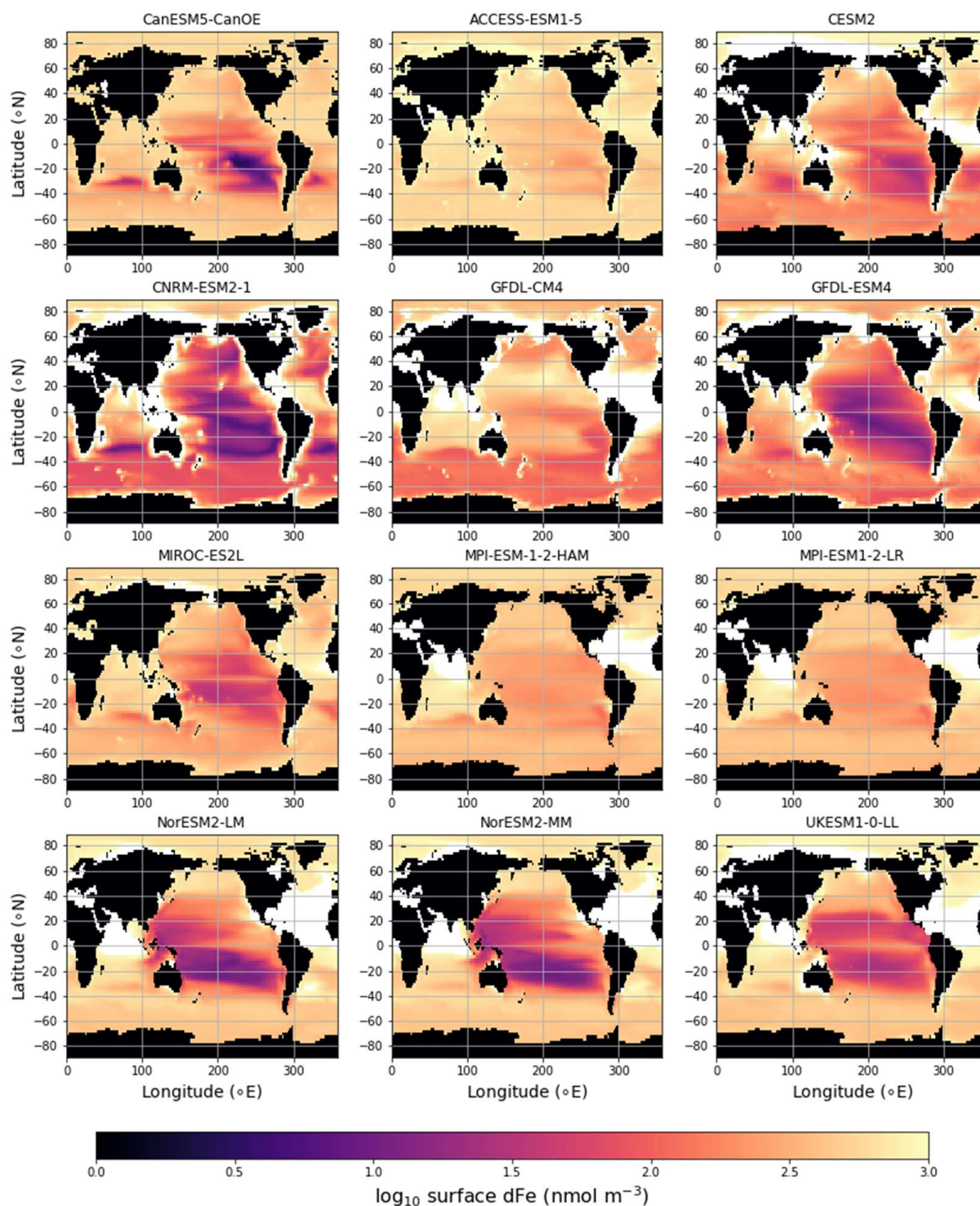


Figure 12 - Global distribution of dissolved iron (dFe) concentration (\log_{10} of concentration in nmol m^{-3}) at the ocean surface for CanESM5-CanOE and other CMIP6 models that published this field. Concentrations exceeding 1000 nmol m^{-3} are masked white.

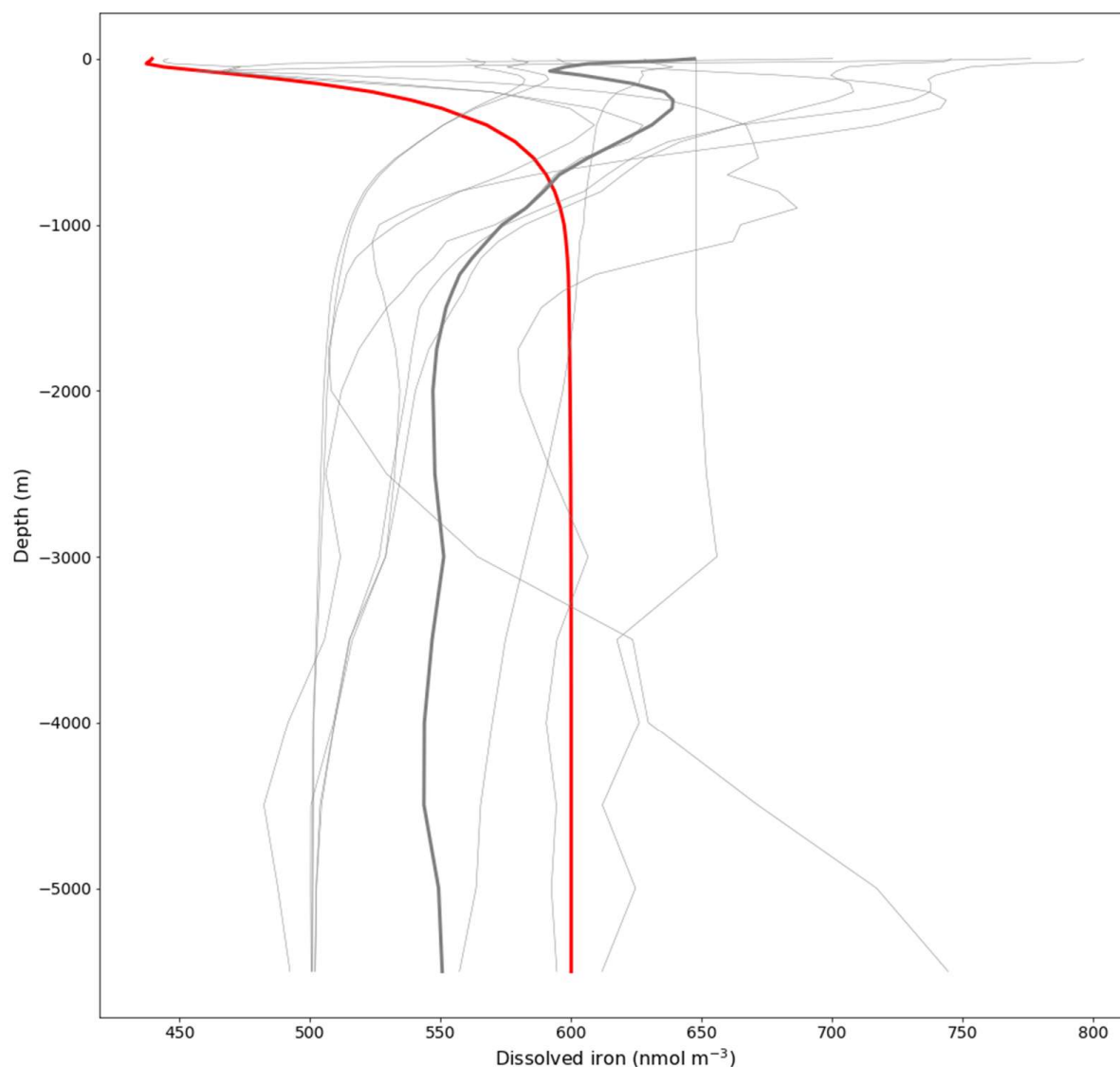


Figure 13 - Global mean depth profiles of dissolved iron concentration for CanESM5-CanOE and other CMIP6 models that published this field. GFDL-CM4 is excluded because it has very high concentrations (>2000 nmol m^{-3}) near the surface. Thick red line represents CanESM5-CanOE, thin grey lines individual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



544 remineralization) profile, some a more “scavenged-type” (maximal at the surface, declining with
545 depth) profile (cf. Li, 1991; Nozaki, 2001), and others a hybrid profile (increasing downward but
546 with a surface enrichment). CanESM5-CanOE is at the “nutrient-type” end of spectrum with a
547 generally monotonic increase with depth to a near-constant deep-water concentration of 0.6 nM
548 and a very slight near-surface enrichment (see also Supplementary Figures S4b,c). In CanESM5-
549 CanOE the scavenging model is very simple, with distinct regimes for concentrations greater or
550 less than 0.6 nM; scavenging rates are very high above this threshold which causes deep-water
551 concentrations to converge on this value. The generally nutrient-like profile suggest that in
552 CanOE the scavenging rate is quite low for concentrations below 0.6 nM. CanOE considers
553 particulate organic matter (POM) as an index of all particulate matter available for scavenging
554 onto, and model POM concentrations fall off rapidly below the euphotic zone (not shown).

555

556 Mean surface nitrate and dFe concentrations for selected ocean regions are shown in Figure 14.
557 CanESM5-CanOE shows concentrations that are within the range of CMIP6 models, although in
558 some cases at the higher or lower end. Surface nitrate concentrations generally compare
559 favourably with the observation-based climatology, but are biased low in HNLC regions other
560 than the Southern Ocean. These biases are not necessarily a consequence of having too much or
561 too little iron. For example, in the Southern Ocean CanESM5-CanOE has among the highest
562 surface nitrate concentrations, but it also has some of the highest dFe concentrations, and the
563 high nitrate bias is present in CanESM5 as well. Comparisons with the limited GEOTRACES
564 data available suggest that near surface dFe concentrations in the Southern Ocean are biased high
565 rather than low in CanESM5-CanOE (not shown). One region where there does seem to be a
566 strong correlation between surface nitrate and dFe concentrations is the western subarctic

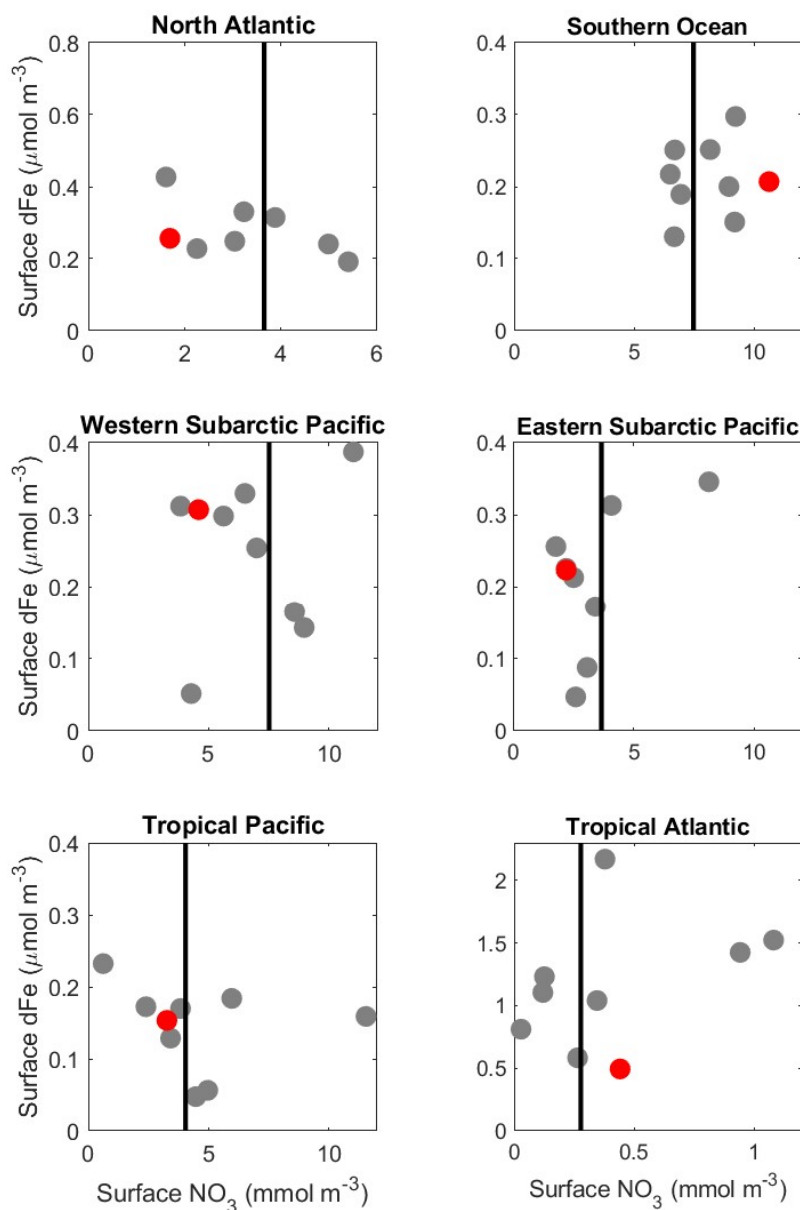


Figure 14 - Mean surface nitrate (NO_3) vs. dissolved iron (dFe) concentrations in different oceans, including the major high nutrient / low chlorophyll (HNLC) regions. CanESM5-CanOE is shown as a red dot and other CMIP5 models as grey dots (CanESM5 is not included because it does not have iron). Observed NO_3 is shown as a vertical black line as there are no observational estimates of dFe concentration. Region definitions are given in Supplementary Table S4.



568 Pacific. All but two models fall along a spectrum from high Fe / low nitrate to low Fe / high
569 nitrate. CanESM5-CanOE falls near the high Fe / low nitrate end of the range.
570
571 Surface nitrate concentrations along the Pacific equator during the upwelling season (June-
572 October) for CanESM5 and CanESM5-CanOE are shown in Figure 15. CanESM5-CanOE better
573 represents the east-west gradient, while CanESM5 has slightly higher concentrations in the core
574 upwelling region. Both models underestimate the highest concentrations around 100°W. Some
575 localized maxima in this data product are due to undersampling; however, examination of
576 ancillary data sets such as satellite sea surface temperature suggests that the enrichment at
577 100°W accurately reflects ocean upwelling (not shown). Although CanESM5 iron limitation is
578 calculated from an earlier version of the same data product, the Fe mask is based on the
579 minimum nitrate concentration over the annual cycle, whereas the data shown here are for the
580 upwelling season. In CanESM5-CanOE, the distribution of surface nitrate is an emergent
581 property of the model, and the fidelity to the observed distribution is generally good.

582

583 **3.4 Plankton biomass, detritus, and particle flux**

584

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

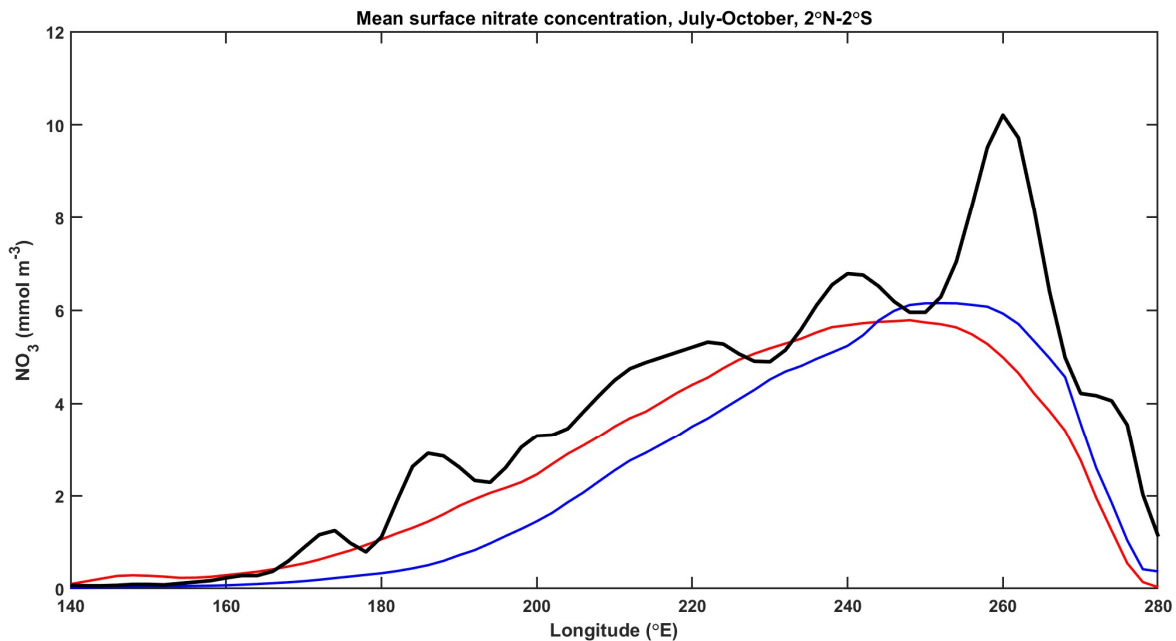


Figure 15 - Surface nitrate (NO_3) concentrations along the Pacific equator (mean from 2°S - 2°N) during the upwelling season (June-October) for CanESM5-CanOE (red), CanESM5 (blue), and WOA2013 observations (black).

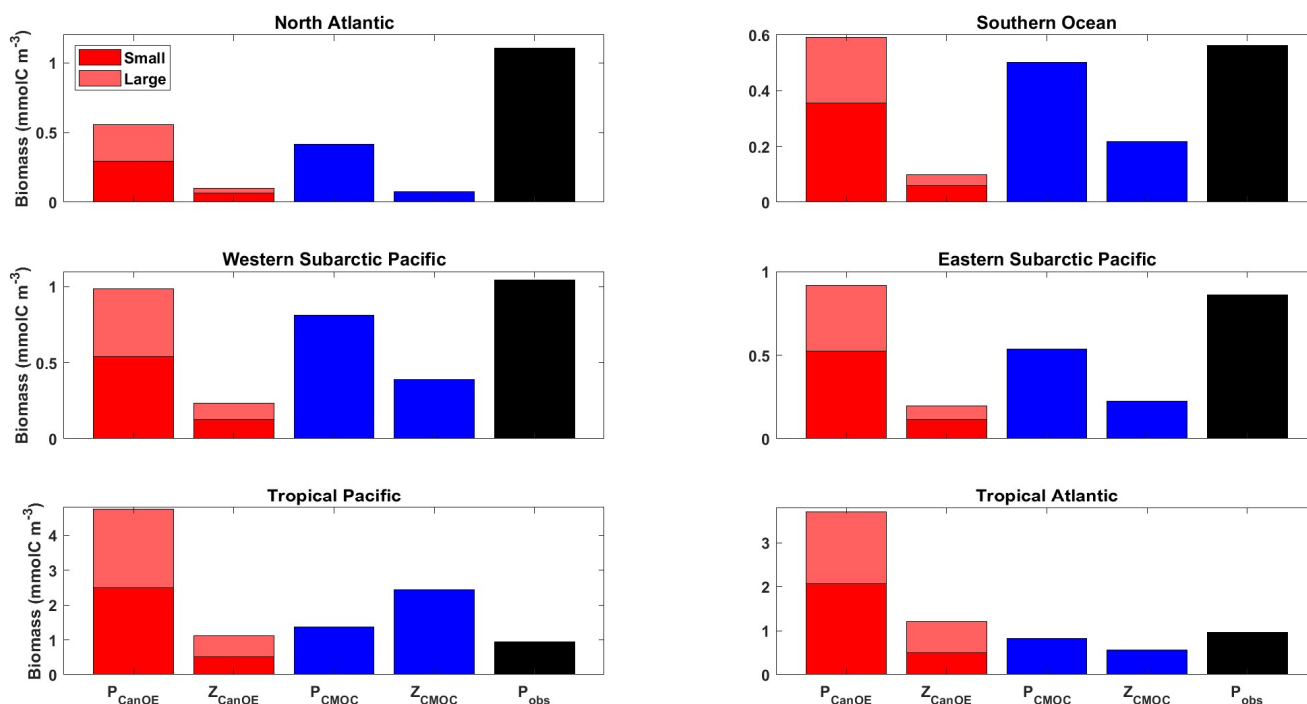


Figure 16 - Annual mean surface ocean concentration of large and small phytoplankton and zooplankton in CanESM5-CanOE (red) and of phytoplankton and zooplankton in CanESM5 (blue) for the representative ocean regions shown in Figure 14. Observational estimates (black) are for phytoplankton biomass calculated 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 S4.



593 phytoplankton have similar abundance, and are substantially more abundant than zooplankton.
594 Part of the rationale for multiple food chains is that they better represent the way that actual
595 plankton communities adapt to different physical ocean regimes and therefore are better able to
596 simulate distinct ocean regions with a single parameter set (e.g., Chisholm, 1992; Armstrong,
597 1994; Landry et al., 1997; Friedrichs et al., 2007). The expectation is that small phytoplankton
598 will be more temporally stable and large phytoplankton will fluctuate more strongly between
599 high and low abundances.

600

601 The mean annual cycles of surface chlorophyll largely conform to the expected pattern, e.g., in
602 the North Atlantic and the western subarctic Pacific large phytoplankton are dominant in summer
603 and much more variable over the seasons (Figure 17). Compared to observations, CanESM5
604 models underestimate the amplitude of the seasonal cycle in the North Atlantic and overestimate
605 it in the North Pacific. CanESM5 shows a stronger and earlier North Atlantic spring bloom
606 compared to CanESM5-CanOE; the observations are in between the two in terms of timing, and
607 both models underestimate the amplitude (Figure 17). In the tropics, the seasonal cycle is weak.
608 The tropical Atlantic shows the expected seasonal cycle but not the expected dominance of large
609 phytoplankton in summer. These size-fractionation patterns are difficult to validate against
610 observations. CanESM5-CanOE generally overestimates the total near surface chlorophyll in
611 both the tropical Pacific and the tropical Atlantic.

612

613 Zooplankton biomass (especially microzooplankton) is also somewhat difficult to test against
614 observations, but our model concentrations appear to be biased low. White et al. (1995), for
615 example, show a transect of vertically resolved mesozooplankton abundance along 140°W in the

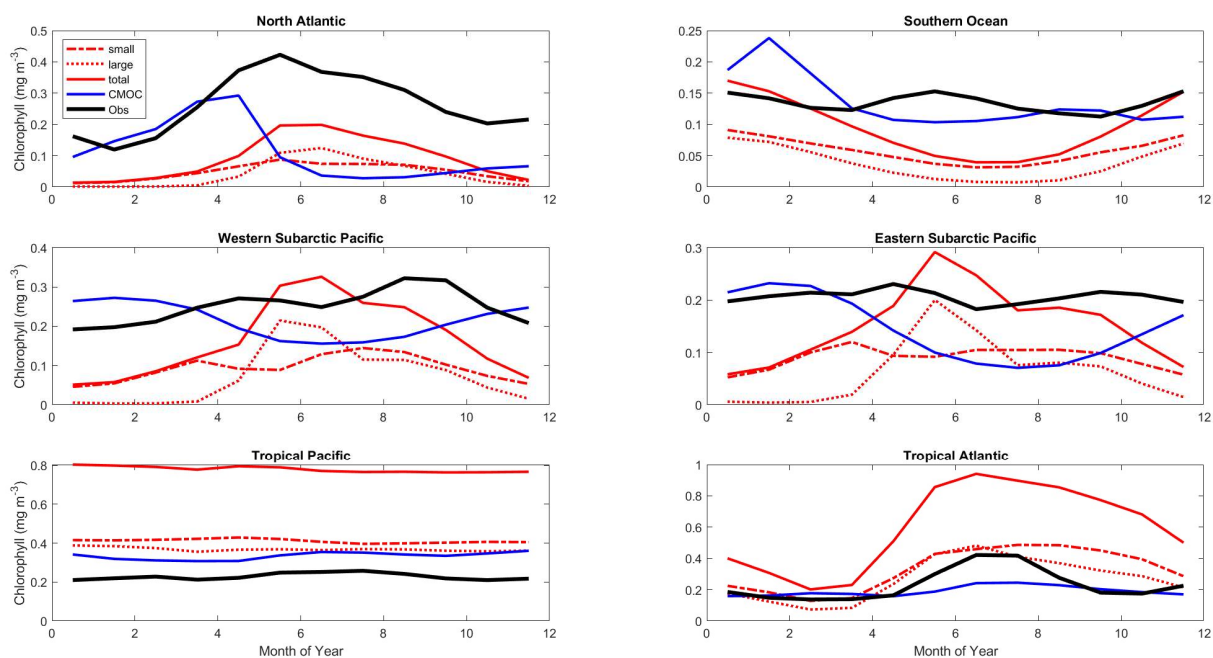


Figure 17 - Mean annual cycle of surface chlorophyll for the representative ocean regions shown in Figures 14 and 16. CanESM5-CanOE large and small phytoplankton concentrations are shown separately and combined (red) along with CanESM5 (blue) and observational estimates (black). Region definitions are shown in Supplementary Table S4.



617 tropical Pacific, with biomass ranging from about 0.1-0.7 mmolC m³; CanESM5-CanOE
618 concentrations in this region are much lower. Stock et al. (2014) estimated depth-integrated
619 biomass of phytoplankton, mesozooplankton, and microzooplankton for a range of oceanic
620 locations in which intensive field campaigns have occurred (estimates of microzooplankton
621 biomass are relatively sparse). They found that in most locations phytoplankton and (combined)
622 zooplankton biomass are of comparable magnitude, whereas in CanESM5-CanOE zooplankton
623 biomass is consistently lower (Figure 16). The global integral biomass of mesozooplankton is
624 about an order of magnitude less than the 0.19 PgC estimated by Moriarty and O'Brien (2013).
625 The CanESM5 total of 0.14 Pg is relatively close to the Moriarty estimate but implicitly includes
626 microzooplankton.

627

628 Surface chlorophyll and POC for CanESM5-CanOE and for ocean colour observational data are
629 shown in Figure 18 (POC in the model is the sum of phytoplankton, microzooplankton, and
630 detrital carbon). The observations have a lower limit for POC that is not present in the model
631 (~17 mgC m⁻³), which is unsurprising given the processes neglected in the model, i.e., in regions
632 of very low chlorophyll there is still substantial dissolved organic carbon, bacteria that consume
633 it, and microzooplankton that consume the bacteria and produce particulate detritus. The
634 observational data show a fairly linear relationship at low concentrations, but with a curvature
635 that implies a greater phytoplankton fraction in more eutrophic environments (cf. Chisholm,
636 1992). The model, by contrast, shows a fairly linear relationship over the whole range of
637 concentrations. In other words, the phytoplankton share of POC is higher and more constant in
638 the model than in the observations. The living biomass (phytoplankton + microzooplankton)
639 fraction of total POC in CanOE is generally in excess of 50% (not shown), which is implausible

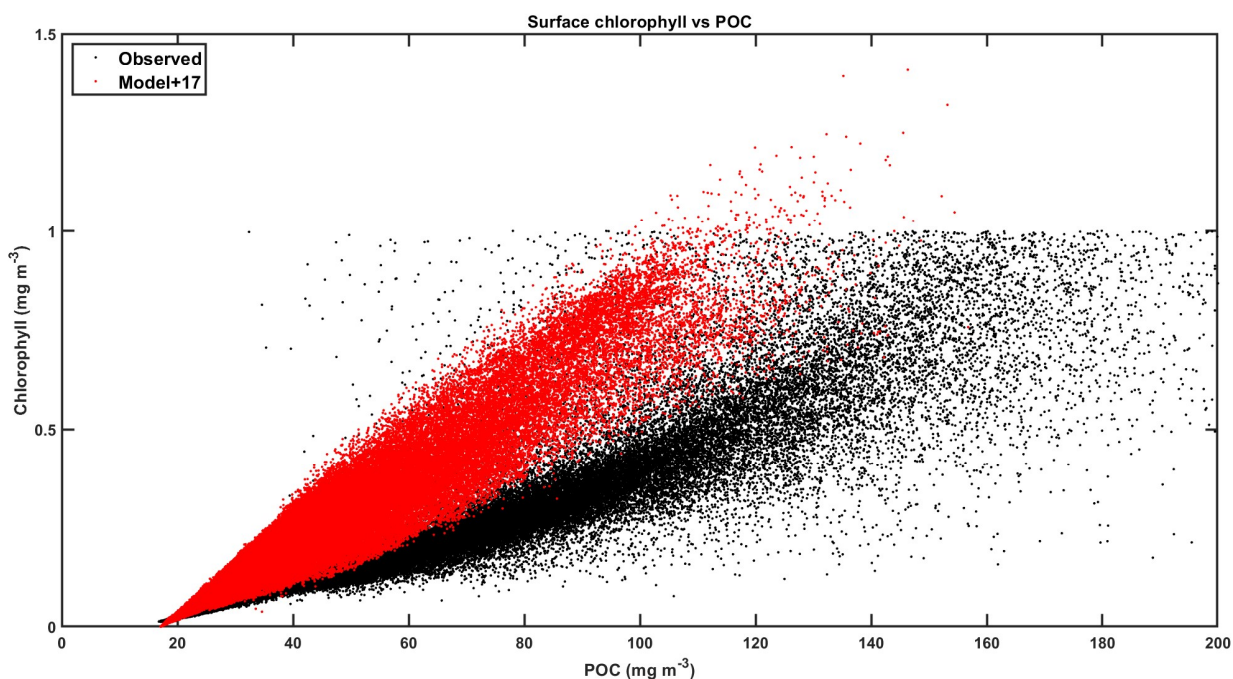


Figure 18 - Climatological surface particulate organic carbon (POC) vs. chlorophyll for CanESM5-CanOE (red) and observations (black). Data are for all ocean grid points ($2 \times 2^\circ$ uniform global grid) for all months of the year where observational data are available. Model POC is offset 17 mg m^{-3} for illustrative purposes.



641 for a real-world oceanic microbial community (e.g., Christian and Karl, 1994) but consistent
642 with the relatively low rates of export from the euphotic zone.

643

644 Export production for a range of CMIP6 models is shown in Figure 19a. CanESM5-CanOE is at
645 the low end of the range. Observations are not shown because the range of observational
646 estimates covers almost the entire range of the plot (e.g., Siegel et al., 2016). Note also that
647 CanESM5 export is quite a bit lower than in CanESM2, which is relatively high for CMIP5
648 models (not shown). The difference between CanESM2 and CanESM5 is attributable primarily
649 to different circulation, although the different initialization fields for nitrate might also play a
650 small role. The lower rate in CanESM5-CanOE is consistent with the above results regarding
651 plankton community structure (e.g., the concentration of detritus is generally low compared to
652 living biomass), as well as the lower sinking rate for small detritus. The latitudinal distribution of
653 export is shown in Figure 19b. CanESM5 shows very high export in the mid-latitudes of the
654 Southern Ocean, similar to CanESM2 (not shown). Both CanESM5 models show latitudinal
655 patterns consistent with the range of other models. CanESM5 has slightly greater export in the
656 equatorial zone; in both CanESM models the equatorial enrichment attenuates very rapidly with
657 latitude and the rates are low in the subtropics.

658

659 **3.5 Historical trends**

660

661 Cumulative ocean uptake of CO₂ is shown in Figure 20 for the historical experiment (1850-
662 2014). CanESM models are biased low relative to observation based estimates (~145 PgC, see
663 Friedlingstein et al., 2020) and the ensemble mean of other models (144 PgC, Figure 20), but fall

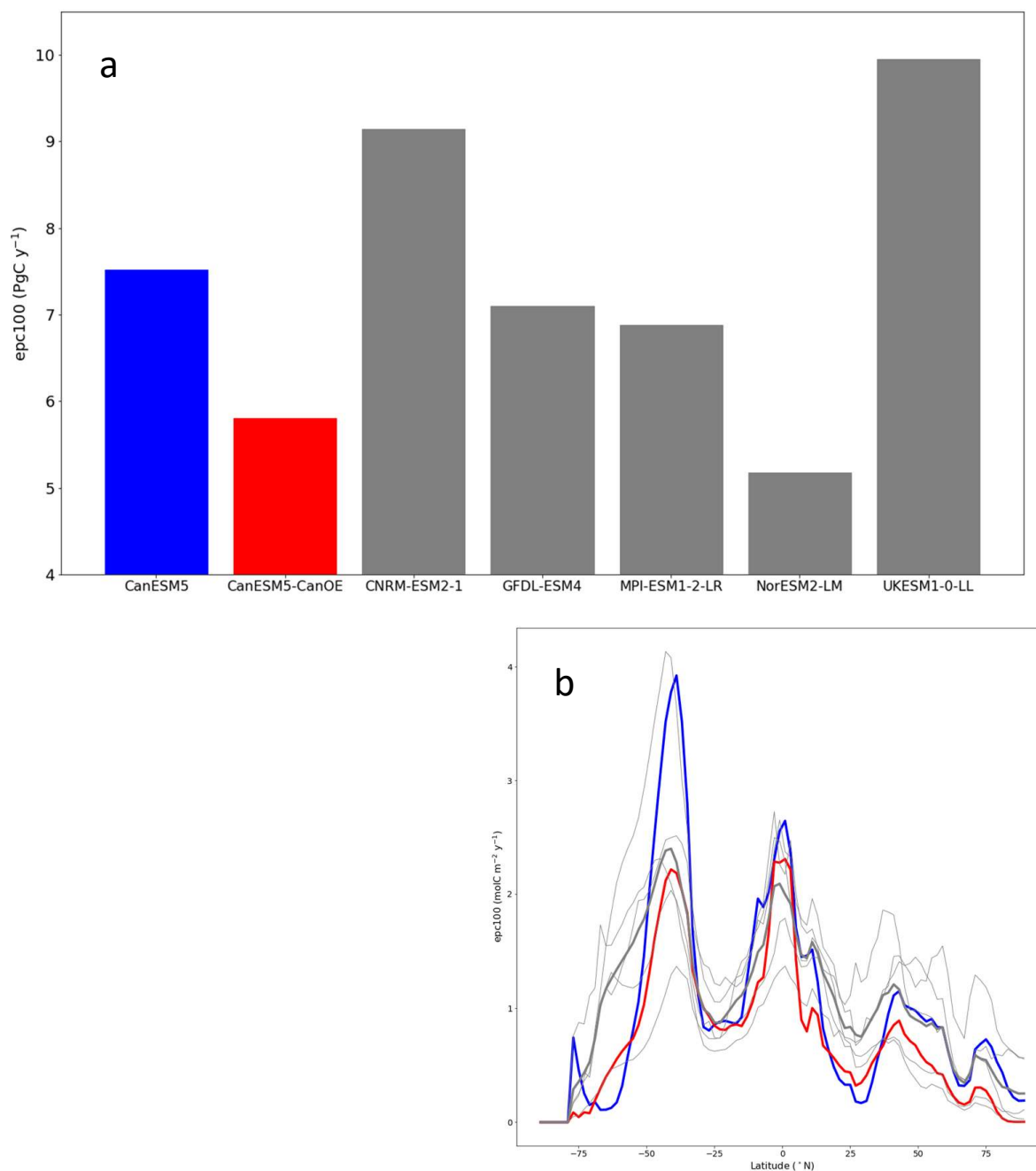


Figure 19 - (a) Global total export production (epc100) in PgC y⁻¹ (b) and zonal mean export production in molC m⁻² y⁻¹ according to selected CMIP6 models (mean for 1985-2014 of historical experiment). Thick red line represents CanESM5-CanOE, thick blue line CanESM5, thin grey lines individual CMIP6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

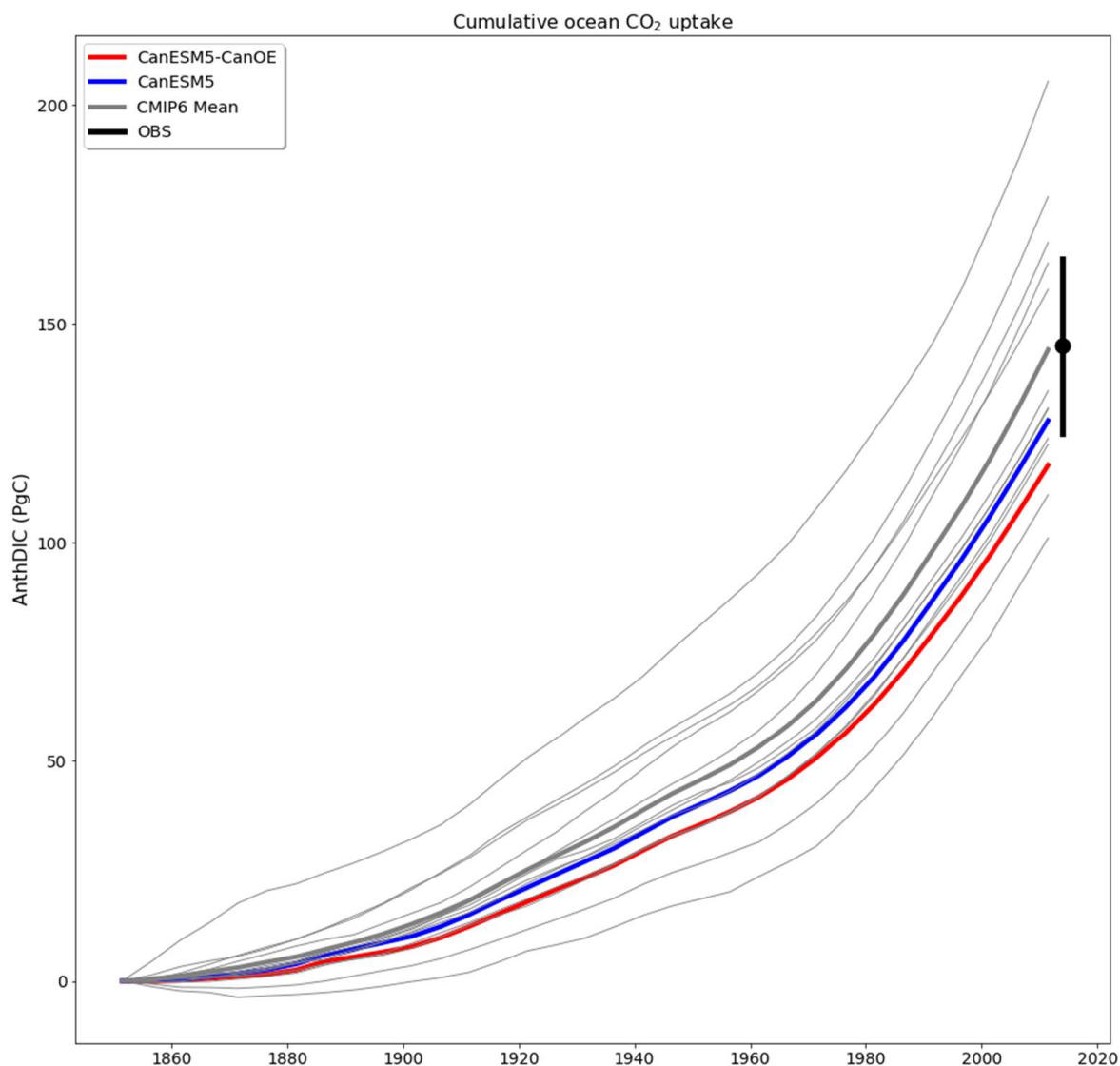


Figure 20 - Cumulative ocean uptake of carbon dioxide (CO_2) as anthropogenic dissolved inorganic carbon (AnthDIC) in PgC over the course of the historical experiment (1850-2014). Data are shown as successive five-year means. CMIP6 mean (thick grey line) indicates ensemble mean for CMIP6 models (thin 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).



666 well within the spread of CMIP6 models. CanESM5-CanOE has lower cumulative uptake than
667 CanESM5 by ~ 10 PgC. As the models were not fully equilibrated when the historical run was
668 launched, this difference does not necessarily arise from the biogeochemical model structure;
669 part of the difference can be attributed to differences in the spinup protocol (cf. Séférian et al.,
670 2016). The drift in the piControl experiment over the 165 years from the branching off of the
671 historical experiment is -10.5 PgC in CanESM5-CanOE and -5.9 PgC in CanESM5, so drift
672 accounts for about half (44%) of the difference in net ocean CO₂ uptake. The vertical distribution
673 of anthropogenic DIC is very similar between CanESM5 and CanESM5-CanOE (not shown).

674

675 The long-term trend in global total export production is shown in Figure 21. The model values
676 must be normalized in order to compare trends, since the differences among means are large
677 compared to the changes over the historical period (Figure 19). CanESM5 shows a greater
678 decline than most other CMIP6 models, while CanESM5-CanOE is more similar to non-
679 CanESM models. Such trends are difficult or impossible to meaningfully constrain with
680 observations, but the general expectation has been that export will decline somewhat due to
681 increasing stratification (e.g., Steinacher et al., 2010). The change in CanESM5 is geographically
682 widespread and not concentrated in a specific region or regions: export is maximal in the tropics
683 and the northern and southern mid-latitudes (Figure 19b) and declines over the historical period
684 in all of these regions (Supplementary Figure S1). In CanESM5-CanOE, export declines in the
685 same regions, but the magnitude of the change is smaller, and in the Southern Ocean increases
686 and decreases in different latitude bands largely offset each other.

687

688

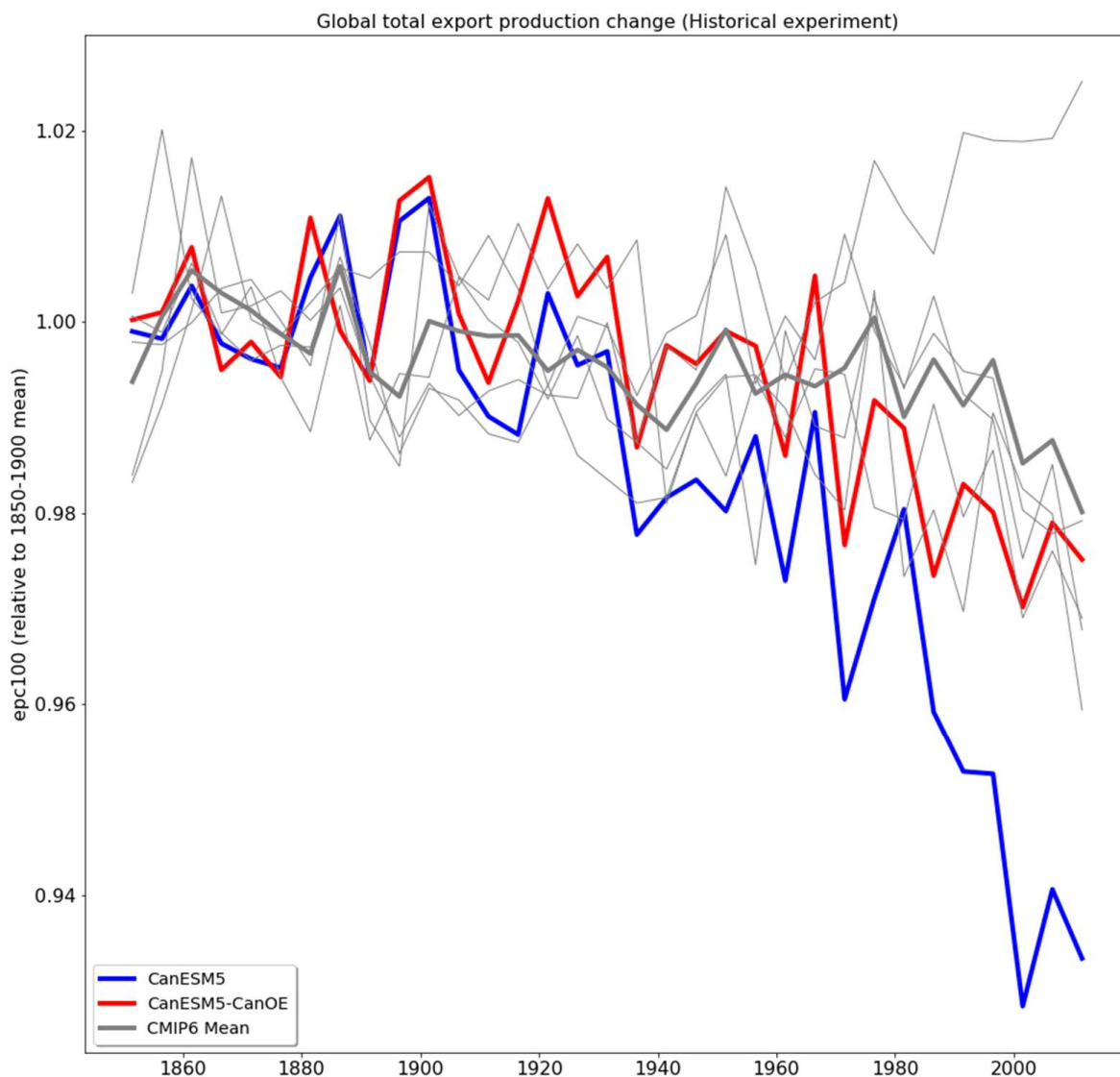


Figure 21 - Change in export production (epc100) over the course of the historical experiment (1850-2014), normalized to the 1850-1900 mean. Data are shown as successive five-year means. Thick red line represents CanESM5-CanOE, thick blue line CanESM5, thin grey lines other CMIP6 models, and thick grey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



690 The trend in the volume of ocean water with O₂ concentration less than 6 or 60 mmol m⁻³ is
691 shown in Figure 22. Again, the totals are normalized to a value close to the preindustrial, as the
692 differences among models are large (Figure 5). For the volume with <60 mmol m⁻³, CanESM
693 models show relatively little change; in CanESM5 the volume actually declines slightly, while in
694 CanESM5-CanOE it increases, but the total change is <1% in each case. As with the baseline
695 volumes, the range among models is large, with one model showing an increase approaching
696 10% of the total volume estimated for WOA2013 (Figures 5b and 22b). For the volume with <6
697 mmol m⁻³ (Figure 22a), CanESM models are among the most stable over time. In CanESM5, the
698 volume again declines, although this is within the range of internal variability. Again some
699 models show fairly large excursions, but in this case none shows a strong secular trend over the
700 last half-century.

701

702 **4. Discussion**

703

704 The development of CanOE was undertaken in response to some of the most severe limitations
705 of CanESM1/2, and in light of our collective experience. In addition to CMOC (Zahariev et al.,
706 2008), previous models developed by members of our group include Denman and Peña (1999;
707 2002), Christian et al. (2002a; 2002b), Christian (2005), and Denman et al. (2006). Christian et
708 al. (2002a) had a prognostic Fe cycle and multiple phytoplankton and zooplankton species, but
709 had fixed elemental ratios. Christian (2005) incorporated a cellular-regulation model, but only
710 for a single species and without Fe limitation. Christian (2005) had prognostic chlorophyll
711 whereas Denman and Pena (1999; 2002) and Christian et al. (2002a) used an irradiance-depend

712

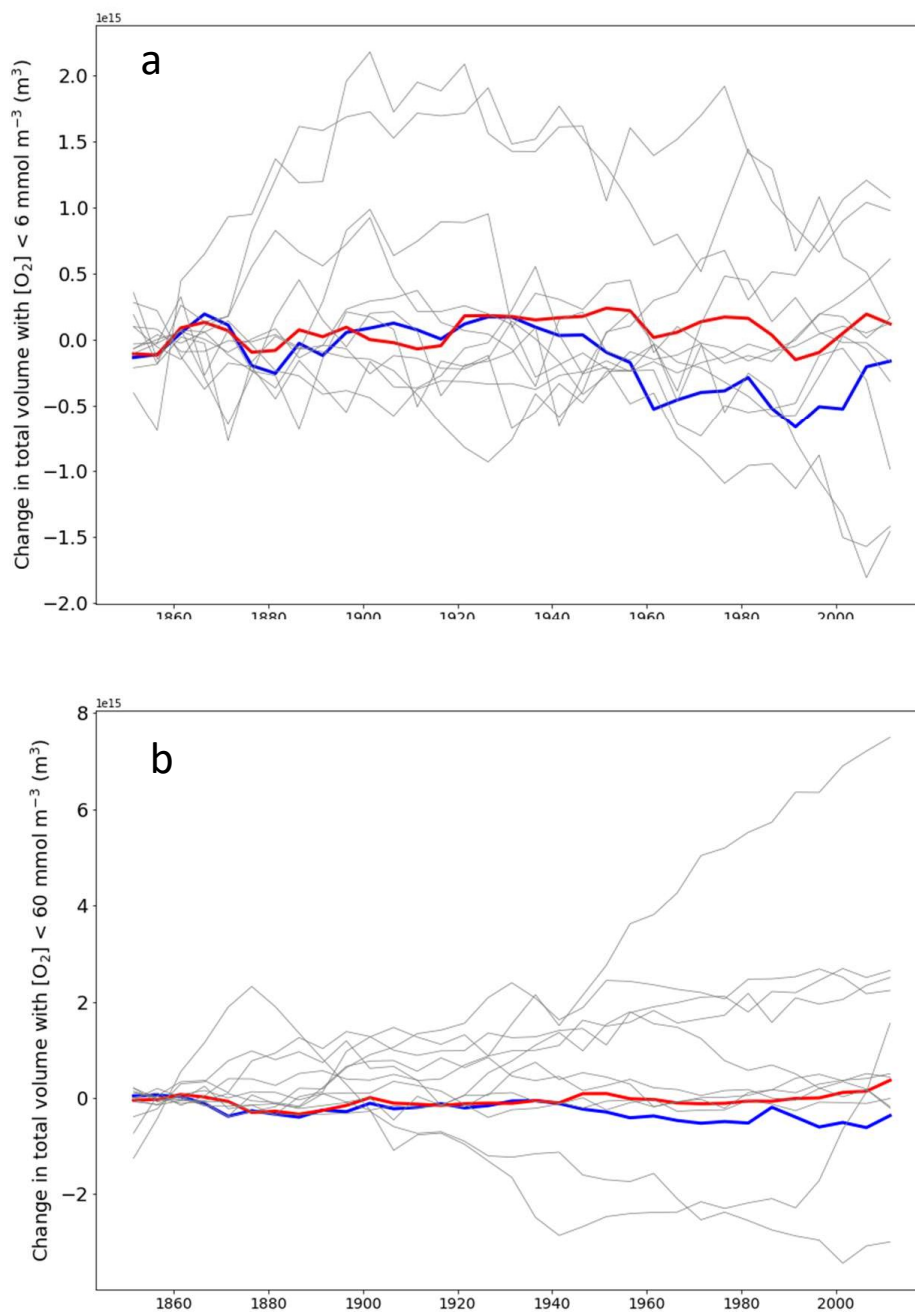


Figure 22 - (a) Change in total ocean volume with oxygen (O_2) concentration less than (a) 6 mmol m^{-3} and (b) 60 mmol m^{-3} over the course of the historical experiment (1850-2014), normalized to the 1850-1870 mean. Data are shown as successive five-year means. Thick red line represents CanESM5-CanOE,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



714 ent diagnostic formulation. Christian et al. (2002a) used multiplicative (Franks et al., 1986) graz-
715 ing, which creates stability in predator-prey interactions but severely limits phytoplankton bio-
716 mass accumulation under nutrient-replete conditions.

717

718 One of the most important lessons from Christian et al. (2002a; 2002b) was that when a fixed
719 Fe/N ratio is employed, sensitivity to this parameter is extreme. Because Fe cell quotas are far
720 more variable than N, P, or Si quotas, treating this parameter as constant results in the specified
721 value influencing the overall solution far more than any other parameter. CanESM5-CanOE
722 largely succeeded in creating a prognostic Fe-N limitation model that produces HNLC conditions
723 in the expected regions (Figures 10, 11, 14, 15), although surface nitrate concentration is low rel-
724 ative to observation-based estimates in some cases. External Fe sources and scavenging parame-
725 terizations will be revisited and refined in future versions. We note that the aeolian mineral dust
726 deposition field employed here is derived from the CanESM atmosphere model; these processes
727 are not presently interactive but could be made so in the future.

728

729 A particular issue with CanESM2 was that extremely high concentrations of nitrate occurred un-
730 der the Eastern Boundary Current (EBC) upwelling regions. This error resulted from spreading
731 denitrification out over the ocean basin so that introduction of new fixed N from N₂ fixation
732 would balance denitrification losses within each vertical column, whereas in the real world deni-
733 trification is highly localized in the low oxygen environments under the EBCs. CanESM2 did not
734 include oxygen, but CanESM5 CMOC incorporates oxygen as a ‘downstream’ tracer that does
735 not feed back on other biogeochemical processes. The incorporation of a more process-based de-



736 nitrification parameterization in CanOE is independent of the many other processes that are pre-
737 sent in CanOE but not in CMOC: a CMOC-like model with prognostic denitrification is clearly
738 an option. We chose not to include explicit, oxygen-dependent denitrification in CanESM5 be-
739 cause we wanted to maintain a CMOC-based model as close to the CanESM2 version as possi-
740 ble, and because oxygen would not then be a downstream tracer that does not affect other pro-
741 cesses.

742

743 CanOE for the most part successfully reproduces the overall distribution of major tracers such as
744 nitrate, oxygen, DIC and alkalinity (and dFe, to the extent that its distribution is known). One
745 could argue that the gains made relative to CMOC are incremental. However, it is also important
746 to note that CanOE explicitly simulates important processes that are highly parameterized or
747 specified in CMOC. For example, the maintenance of HNLC regions is hardwired into CMOC
748 by specifying iron limitation as a function of the present-day observed distribution of surface ni-
749 trate. Both models show substantial gains in skill relative to CanESM2. These gains are similar
750 in the two models and in coupled or ocean only (with CanESM2 forcing) mode (not shown) and
751 are, therefore, attributable primarily to improvements in the ocean circulation model, although
752 differences in initialization and spinup may also play a small role (e.g., Séférian et al., 2016).

753

754 Plankton community structure in CanOE is somewhat biased toward high concentrations of phy-
755 toplankton, low concentrations of zooplankton and detritus, and low export (Sect. 3.4). In the de-
756 velopment phase, a fair number of experiments were conducted with various values of the graz-
757 ing rates and detritus sinking speeds. A wide range of values of these parameters was tested, with
758 no resulting improvement in the overall results. Possibly the detrital remineralization rates are



759 too high, although primary production is also on the low end of the CMIP6 range (not shown),
760 and would probably decline further if these rates were decreased. The model was designed
761 around the Armstrong (1994) hypothesis of ‘supplementation’ vs ‘replacement’, i.e., small phy-
762 toplankton and their grazers do not become much more abundant in more nutrient-rich environ-
763 ments but rather stay at about the same level and are joined by larger species that are absent in
764 more oligotrophic conditions (see also Chisholm, 1992; Landry et al., 1997; Friedrichs et al.,
765 2007). The results presented here suggest that this was partially achieved but further improve-
766 ment is possible (Figure 17).

767

768 As to whether the gains in skill with CanOE justify the extra computational cost, Taylor dia-
769 grams (Figures 4, 8, 9, and Supplementary Figure S2) show a modest but consistent gain across
770 variables and depths, especially for alkalinity at mid-depths (Supplementary Figure S2), for
771 which CanEM5 displays the least skill relative to other fields or depths. Other processes that are
772 highly parameterized in CMOC, such as calcification and CaCO_3 dissolution, were not addressed
773 in detail in this paper, but are an important factor in determining the subsurface distribution of
774 alkalinity. As noted above for maintenance of HNLC conditions, we emphasize that we are simu-
775 lating as an emergent property something that is parameterized in CMOC, and doing at least as
776 well in terms of model skill. As a general rule, the potential for improving skill and achieving
777 better results in novel environments (e.g., topographically complex regional domains like the
778 Arctic Ocean and the boreal marginal seas), is expected to be greater in less parameterized mod-
779 els (e.g., Friedrichs et al., 2007; Tesdal et al., 2016).

780



781 An updated version of CMOC with prognostic denitrification is clearly possible. However, for
782 the reasons discussed above, a prognostic Fe cycle with a fixed phytoplankton Fe/N remains
783 problematic, and the model would still have a single detritus sinking speed and remineralization
784 length scale. We are also developing CanOE for regional downscaling applications (Hayashida,
785 2018; Holdsworth et al., 2021), and it is likely that the simplification of having a single particle
786 sinking speed is not well suited to a domain with complex topography and prominent continental
787 shelf and slope. The number of tracers in CanOE is not particularly large compared with other
788 CMIP6 models. We expect to further refine CanOE and its parameterizations, evaluate it against
789 new and emerging ocean data sets (e.g., GEOTRACES, biogeochemical ARGO), and incremen-
790 tally improve CMOC (which we will maintain for a wide suite of physical-climate experiments
791 for which ocean biogeochemistry is not central to the purpose). For CMIP6, we chose to keep
792 CMOC as close to the CanESM2 version as possible. This strategy allows us to quantify how
793 much of the improvement in model skill is due to the physical circulation, which is in fact sub-
794stantial (e.g., Figure 8). The CanESM terrestrial carbon model is also undergoing important new
795 developments (e.g., Asaadi and Arora, 2021) and we expect CanESM to continue to offer a cred-
796 ible contribution to global carbon cycle studies, as well as advancing regional downscaling and
797 impacts science.

798

799 *Code availability.* The full CanESM5 source code is publicly available at
800 gitlab.com/ccma/canesm; within this tree the CMOC/CanOE code can be found at
801 gitlab.com/ccma/cannemo/-/tree/v5.0.3/nemo/CONFIG/CCC_CANCPL_ORCA1_LIM_CMOC
802 or [CCC_CANCPL_ORCA1_LIM_CANOE](https://gitlab.com/ccma/cannemo/-/tree/v5.0.3/nemo/CONFIG/CCC_CANCPL_ORCA1_LIM_CANOE) (last access: 21 September 2021). The version of the
803 code which can be used to produce all the simulations submitted to CMIP6, and described in this



804 paper, is tagged as v5.0.3 and has the associated DOI: <https://doi.org/10.5281/zenodo.3251113>
805 (Swart et al., 2019b).

806

807 *Data availability.* All CanESM5 simulations conducted for CMIP6, including those described in
808 this paper, are publicly available via the Earth System Grid Federation (ESGF). All observational
809 data and other CMIP6 model data used are publicly available.

810

811 *Author contributions.* Formulation of the overall research goals and aims: JRC, KLD, NS, NCS;
812 Implementation and testing of the model code: JRC, HH, AMH, WGL, OGJR, AES, NCS; Car-
813 rying out the experiments: JRC, WGL, OGJR, AES, NCS; Creation of the published work: JRC,
814 HH, AMH, AES, NS, NCS.

815

816 *Competing interests.* The authors declare that they have no conflict of interest.

817

818 *Disclaimer.* CanESM has been customized to run on the ECCC high-performance computer, and
819 a significant fraction of the software infrastructure used to run the model is specific to the indi-
820 vidual machines and architecture. While we publicly provide the code, we cannot provide any
821 support for migrating the model to different machines or architectures.

822

823 *Acknowledgments.* This work was made possible by the combined efforts of the CCCMa model
824 development team and computing support team. We thank the data contributors to and develop-
825 ers of the observation-based data products, the NASA ocean colour team, and all of the CMIP6



826 data contributors. The Python packages mocsy by Jim Orr and SkillMetrics by Peter Rochford
827 were invaluable tools in the analysis. William Merryfield and Andrew Ross made useful com-
828 ments on an earlier draft. This paper is dedicated to the memory of Mr. Fouad Majaess, who sup-
829 ported CCCMa supercomputer users for many years and passed away suddenly in 2020.

830

831

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