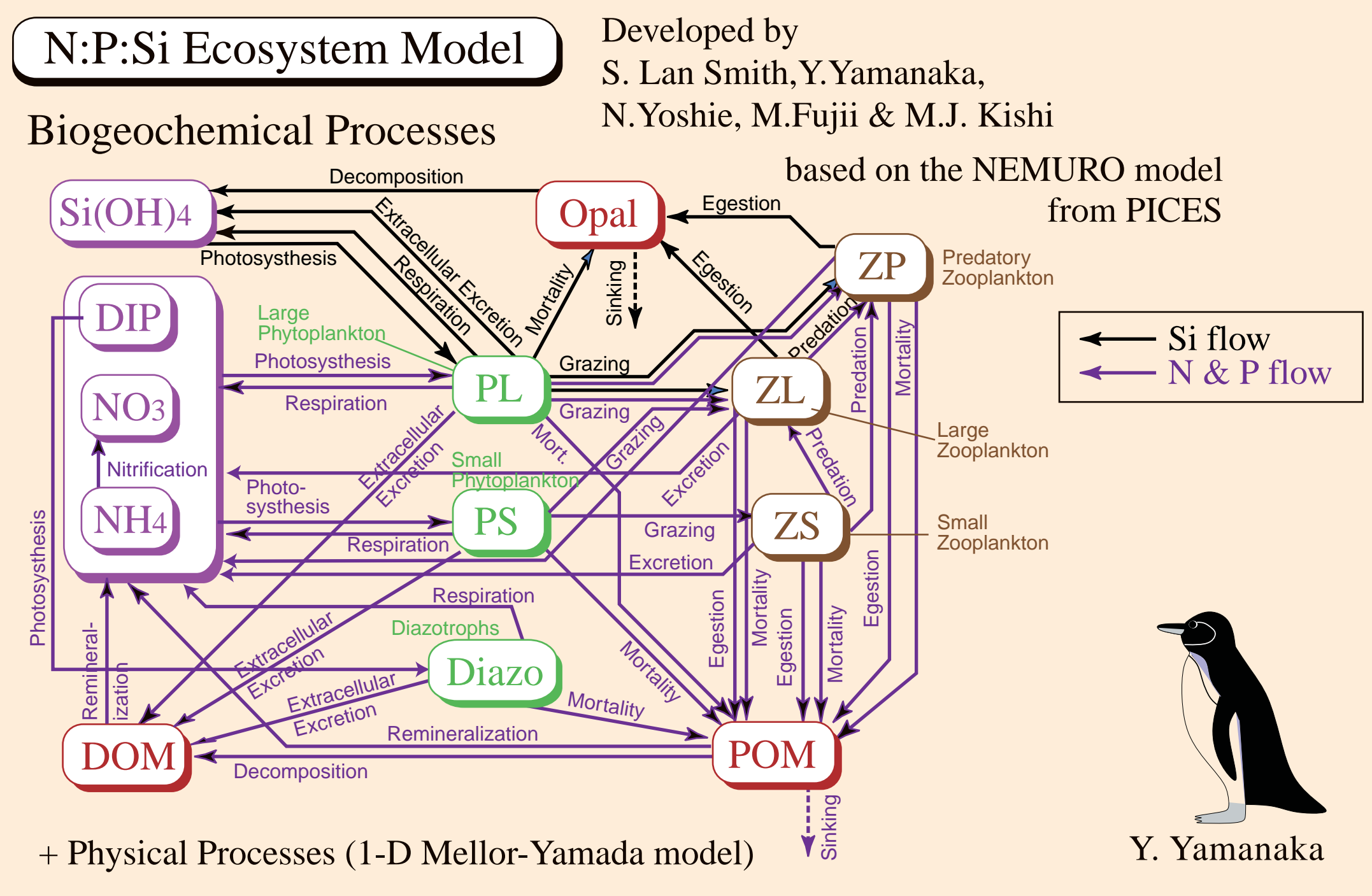


An N-, P- and Si- based Model of Primary Production and Export Applied to Station ALOHA: Can we get the model to agree with the data for primary production, DOM concentrations and POM flux?

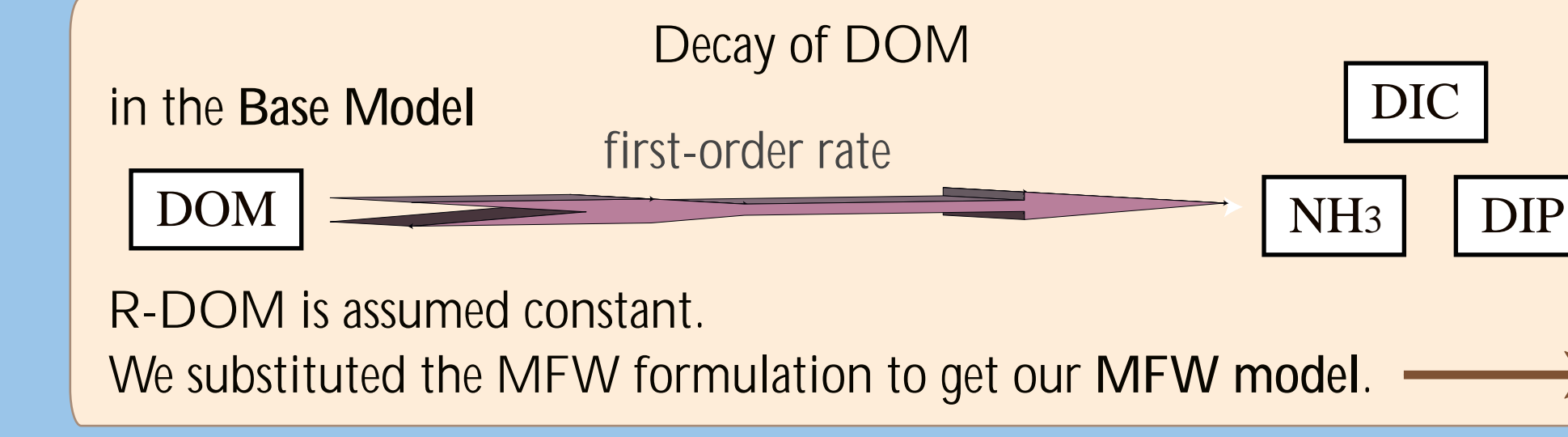
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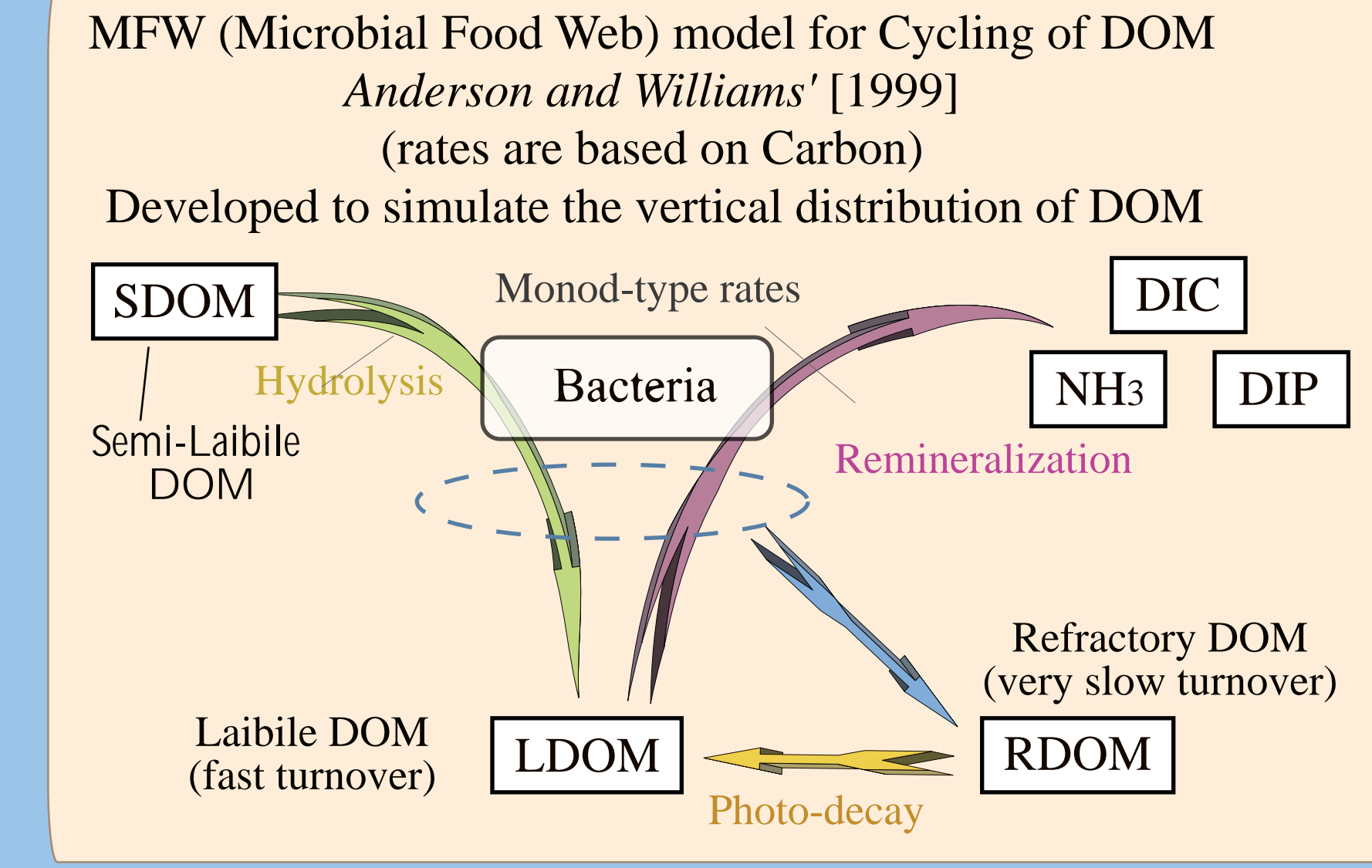
Our modifications to the NEMURO formulation
Diazotrophs (N₂ fixers) based on the model by Fennel et al [2002].
Phosphorous cycles and components for DIP, DOP & POP.
Carbon cycles & components for DIC, DOC & POC
← This is what we call our **Base Model** for this presentation.



Stoichiometries (C:N:P ratios) are fixed for all living organisms & R-DOM.

| component | C:N | N:P | sources |
|-----------|-------|------|---|
| PS, PL | 6.625 | 14* | assumed Redfield; *N:P from Fennel et al |
| Diazo | 6.625 | 45 | Fennel et al [2002] |
| ZS | 5.5 | 14* | Anderson [1992], value for protozoa |
| ZL, ZP | 4.9 | 14* | Anderson [1992], value for copepods |
| Bacteria | 5.1 | 9* | Goldman [1987] |
| RDOM | 20.2 | 54.1 | Averages for all Stn. ALOHA Data from 1997-98 for depths >= 750 m |

Deep Conc. of RDOM: RDOC = 46, RDON = 2.3, RDOP = 0.038 mmoles / liter
Compositions of all other DOM & POM are variable.



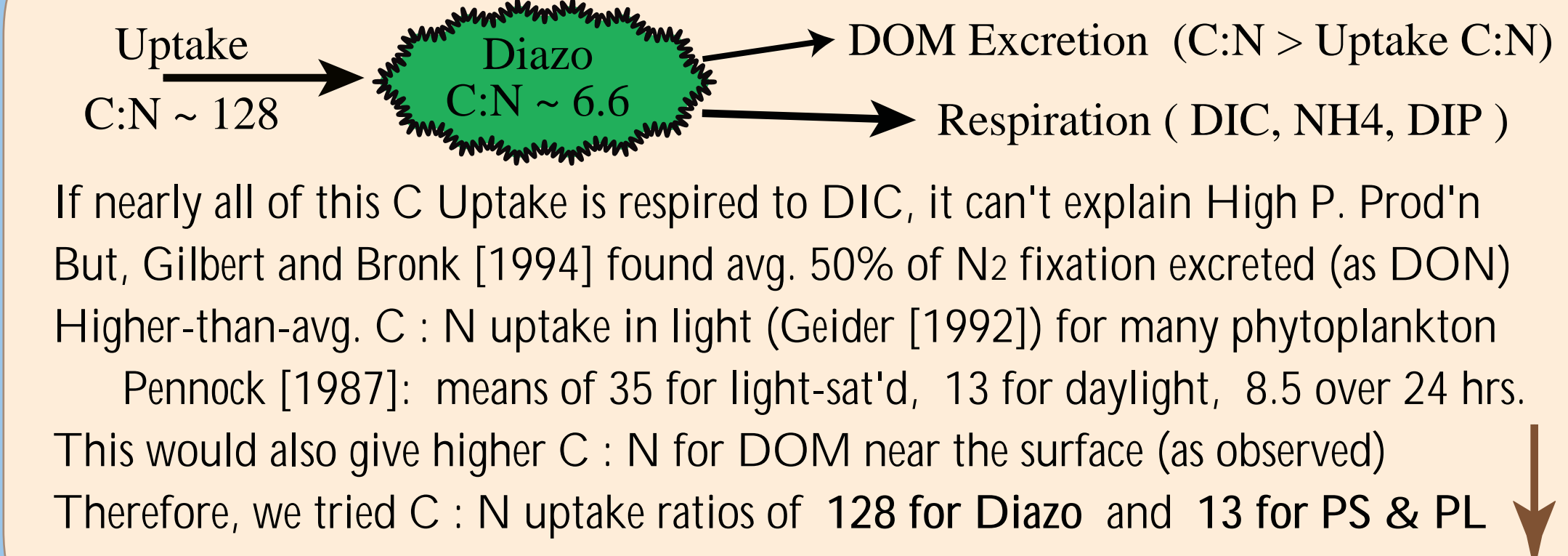
Data for Forcing & Comparison

- HALE-ALOHA Buoy (Pierre Flament, Univ. of Hawaii at Manoa) Wind Velocities & Solar Radiation @ ~ 10 minute intervals
- Cruise Data (approx. monthly) (JGOFS/ HOT-DOGS website) (NO₃ + NO₂), DIP, Si <http://hahana.soest.hawaii.edu/hot/hot-dogs/interface.html>
- DOC, DON, DOP, PN (Particulate Nitrogen), PP (Particulate Phosphorous) Fluxes of POC, PON & POP from Sediment traps

To Drive the Model for 1997 & 1998
30 min. Averages of Solar Radiation, Wind Stress (calc'd from data)
Monthly T & S from cruises, used to damp from 0 to 150 m depth (no data for Jan., 1997; interpolated betw. Dec., 1997 & Feb., 1997)
Minimum diffusive mixing coeff. of 0.5 cm² / s; Downwelling of 5 m / yr (est., Y. Sasai)
Spin-up 1 year, then run: 1997 forcing Once, then 1997 & 1998 simulation

High Uptake Ratios of C:N
Could allow higher Primary Production (C) per mole of N, P
Orcutt et al's [2001] Study of Trichodesmium spp. at BATS measured daytime C : N Uptake mean of 128 (range: 14 - 425) but Biomass remaining near 6.625
They claim this could explain high drawdown of DIC, yet cite studies that found low Excretion (7% of prod., as DOC)

Initial conditions for nutrients and DOM were fixed by interpolation from a spline fit vs. depth for all data from 1997-1998.



Parameters
based loosely on the NEMURO default values for the N. Pacific, but Mortality & Grazing rates were increased to reduce suspended PN, PP concs.
Almost all rates are temperature-dependent, with Q₁₀ = 2
Excretion as a fraction of Gross Production for phytoplankton
Diazo & PS 0.3 (Hood et al, 2002), PL 0.14 (NEMURO std.) for MFW, from Anderson and Williams [1999]
rates were assumed to apply at 20 deg. C, and a Q₁₀ of 2 was applied for each
Degradation of POM is based on measured rates by Fujii et al [2002]
1st order rate: k = 0.15 / day (@ 20 deg. C); Q₁₀ = 2, assuming sinking rate = 6.4 m / day

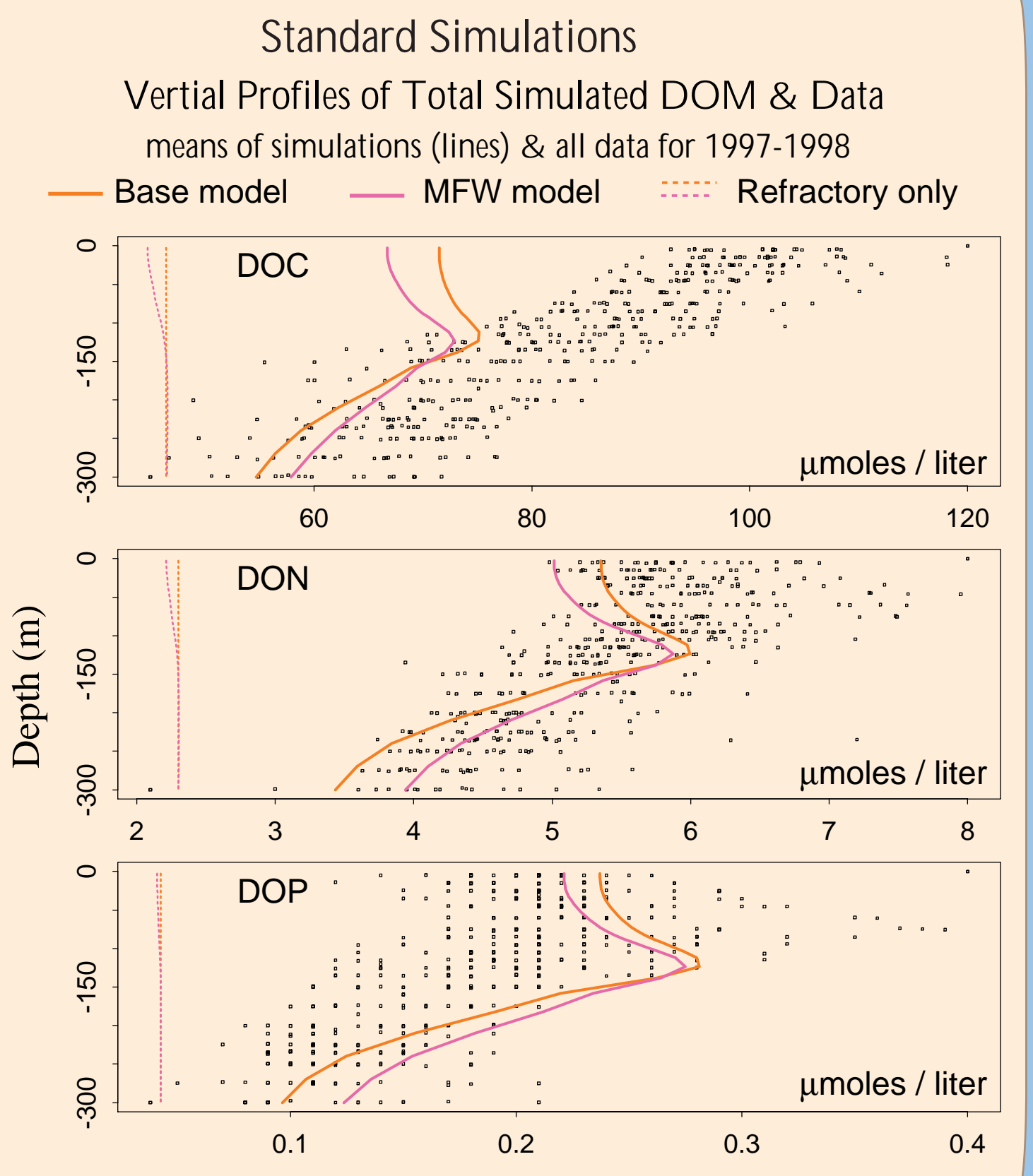
Parameters Tuned for Each Pair of Simulations
to simulate Nutrients, DOM & POM data
Base Model: Decay rate of DOM
MFW Model: Mortality rate of Bacteria

Table 1: Simulations and Data

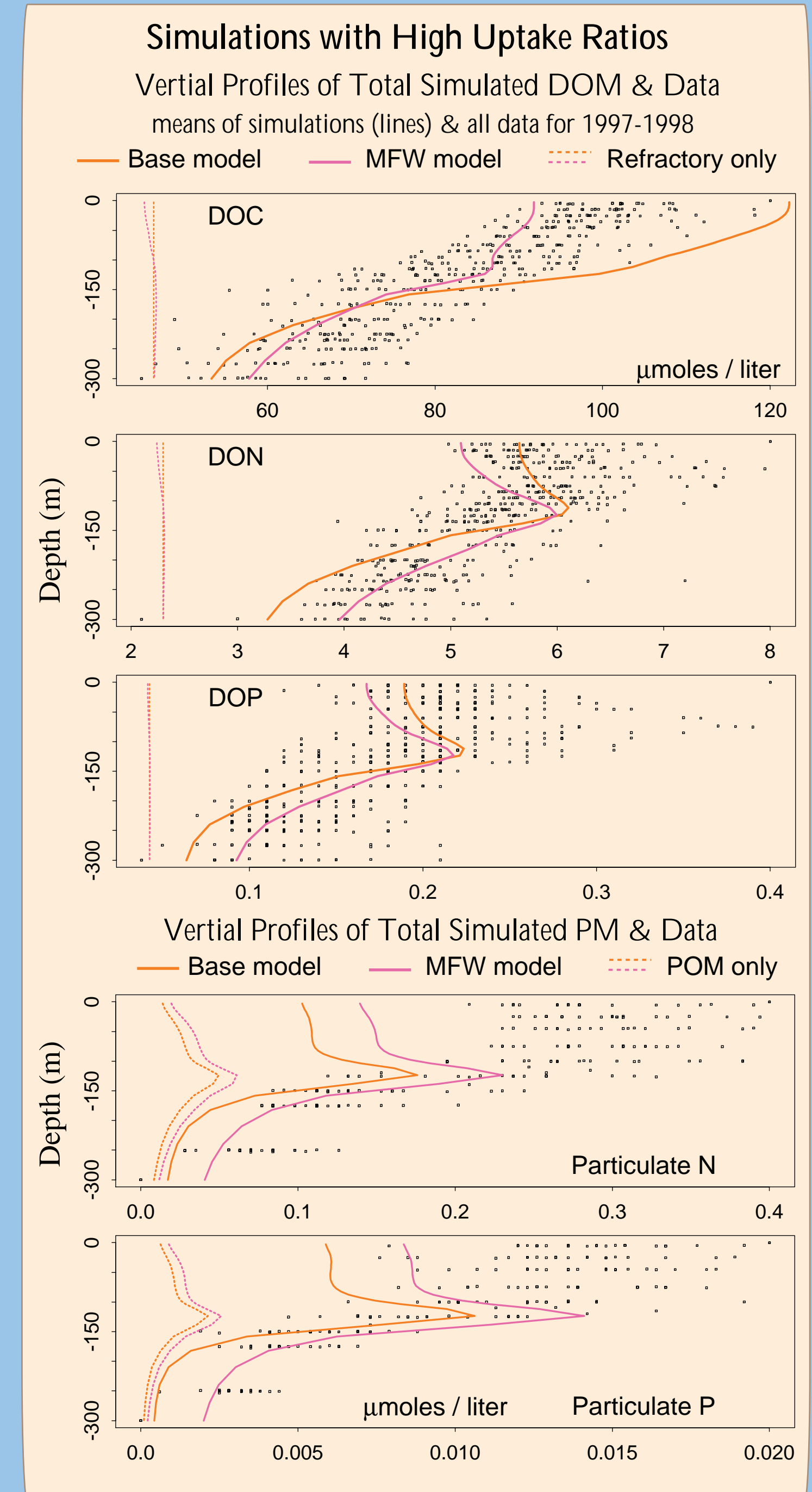
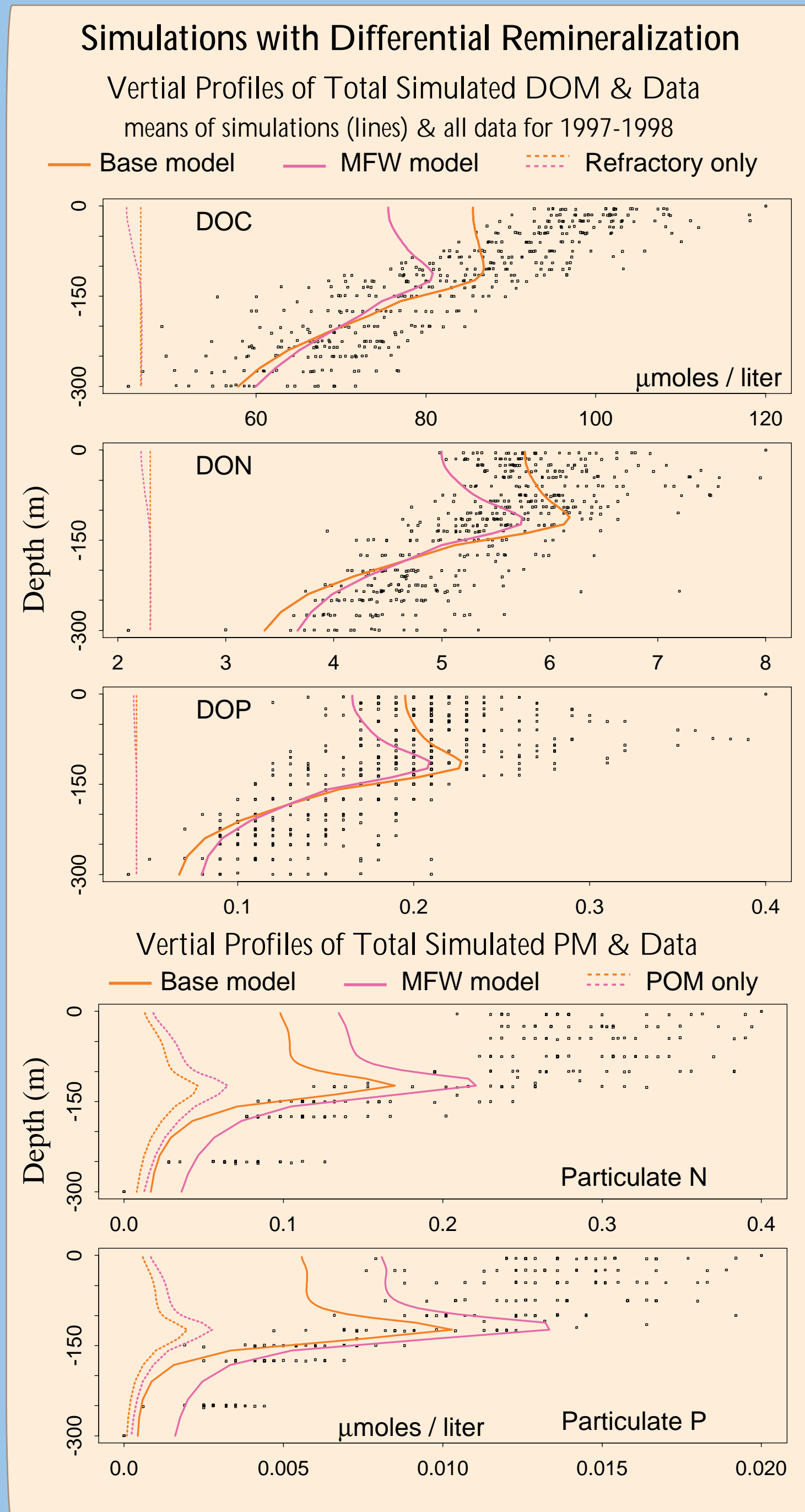
| Quantity (mean for 1997-98) | Data | Standard | | Diff. Remin. | | High Uptake | |
|---|-------|----------|------|--------------|------|-------------|------|
| | | Base | MFW | Base | MFW | Base | MFW |
| Fluxes (mg m ⁻² day ⁻¹), mean of meas. & sim. for all cruises @ 150 m | | | | | | | |
| POC Flux | 30 | 18 | 20 | 28 | 37 | 29 | 35 |
| PON Flux | 4.2 | 3.3 | 3.7 | 3.9 | 5.3 | 4.1 | 5.2 |
| PON Flux | 0.36 | 0.51 | 0.58 | 0.33 | 0.44 | 0.35 | 0.44 |
| Molar Ratios, , mean of meas. & sim. for all cruises @ 150 m | | | | | | | |
| C:N ratio of POM | 8.4 | 6.5 | 6.2 | 8.4 | 8.2 | 8.4 | 7.8 |
| P:N ratio of POM | 26.9 | 14 | 14 | 26 | 27 | 26 | 26 |
| Export of C at 150 m (mg C m ⁻² day ⁻¹), mean for 2 yrs. simulated | | | | | | | |
| as DOC | 13* | 15 | 17 | 20 | 21 | 40 | 31 |
| as POC | 30 | 18 | 20 | 28 | 37 | 30 | 35 |
| Primary Production (mg C m ⁻² day ⁻¹), mean of meas. & sim. for all cruises | | | | | | | |
| Total Primary Production | 478 | 34 | 28 | 69 | 80 | 206 | 162 |
| N ₂ Fixation (mmoles N m ⁻² yr ⁻¹), mean for 2 yrs. simulated | | | | | | | |
| N ₂ Fixation | below | 3 | 3 | 38 | 38 | 36 | 38 |

Data-based estimates of N₂ Fixation range from 31 +/- 18 to 51 +/- 26 mmoles N m⁻² yr⁻¹ (Fennel et al [2002])
* estimate of DOC export based on a spline fit to all data (97-98) and the min. diffusive mixing coefficient

Conclusions
Differential Remineralization (DR) allows the (simulated) ecosystem to retain nutrients near the surface and increase total production by ~100 % for the Base model and by ~200 % for the MFW model (over the Standard Simulations).
DR also improves significantly the simulated profiles of DOC and DOP.
N₂ fixation in the Standard Simulations is limited severely by low DIP concentrations, but is consistent with data-based estimates and other models in all simulations using DR.
The total primary production simulated using the High Uptake C : N ratios is higher by a factor of 2 to 3 than that in the DR simulations with Redfield stoichiometry for uptake (C : N = 6.6). This corresponds to production of large amounts of C-rich DOM which is then degraded relatively slowly (& might not decay during Prim. prod. incubations).
Even so, the simulated primary production values are only ~ 1/3 of the measured primary production.
The simulated DOC is also much better (at least for the MFW model).
The simulations with DR do however seem to overestimate DOC export significantly.
More (better) tuning will almost certainly give better agreement with the data.
(The very high DOC concentrations for the base model in the High Uptake Sims. could be remedied by tuning DR parameters.)
But what can we say from just fitting our model ?
We can test whether ideas like DR and High C : N Uptake can account for the high primary production at places like Stn. ALOHA & BATS, and maybe improve our ability to simulate carbon & nutrient cycles.



In the Standard Simulation, both models simulate DON better than DOC & DOP. Both underestimate DOC and overestimate DOP. Simulated primary production is also only about ~10% of observed.
The C:N & N:P ratios of POM in the simulations are both significantly lower than observed, meaning that the models are exporting too much N and P for each mole of C.
Preferential Remineralization of both POM and DOM is a means by which the ecosystem can retain nutrients in the shallower waters and maintain higher production.
We have introduced parameters $g_{N,P,POM}$ and $g_{N,P,DOM}$ to fix the ratios of the rates of remineralization of C:N and N:P, respectively, for POM. Analogous parameters are applied for degradation of non-refractory DOM (for hydrolysis of SDOM in the MFW model). Setting these parameters to unity results in the standard formulation in which rates follow the stoichiometry of OM.
These are our **Differential Remineralization (DR) Simulations**. The values of the g s are:
 $g_{N,POM} = 0.70$, $g_{P,POM} = 0.50$
 $g_{N,DOM} = 0.60$, $g_{P,DOM} = 0.60$
Thus C is remineralized more slowly than N, which is remineralized more slowly than P.
The fits to DOP & DOC are better, but the concentration and C : N ratio of DOM near the surface are too low.



For the **High Uptake Simulations**, the g s are:
 $g_{N,POM} = 0.70$, $g_{P,POM} = 0.50$
 $g_{N,DOM} = 1.15$, $g_{P,DOM} = 0.60$
Thus DOC is remineralized faster than DON, which is remineralized more slowly than DOP.

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