To the editor:

Please find our point-by-point responses to the reviewer comments below. We have addressed all major comments in the updated manuscript. A latex-diff generated file is also included at the end of this document. Unfortunately, the resulting document produced some errors and does not highlight the changes very well. Therefore, we list the major changes to the manuscript below for your reference:

- Following the suggestion of Reviewer 1, we added a new section describing the timestepping methodology of the model (Section 4). We have also included a new figure demonstrating the capability of the adaptive timestepping scheme (Figure 4), which is discussed in the model validation section.
- To address Reviewer 1's comments on the Antarctica simulations, more information has been provided on the similations in the Antarctica section, including a new figure showing the error in ice thickness and a scatterplot of modeled versus observed surface velocity (Figure 11).
- Following the suggestion of Ed Bueler, we have included a new figure showing Yelmo's performance for the Halfar dome benchmark test (Figure 3), which is discussed in the model validation section.
- To address Ed Bueler's concerns regarding the thermodynamics solver, we have included two new figures that show Yelmo's performance for the enthalpy benchmark experiments of Kleiner et al. (2015) (Figs. 5 & 6), which are discussed in the model validation section.
- Since first submitting the manuscript, we have included the local evolution of the basal water layer as a prognostic variable in the model, including a new equation (Eq. 29).

These changes, as well as all additional modifications, are discussed below. Thank you for your consideration.

	Short comment 1	(Matthew Hoffman))
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Panel b of Figure 2 is very similar to panel b of Figure 1 in Hoffman, et al. (GMD, 2018, https://www.geosci-model-dev.net/11/3747/2018/gmd-11-3747-2018.html), see attached image. The authors may want to consider referencing the previous paper in the figure caption.

Thank you very much for the comment. Indeed Fig. 1 from Hoffman et al. (2018) inspired our own figure. We are happy to make the reference more explicit, and have done so in the caption of the figure revised manuscript.

	Reviewer	comment 1	(anony	ymous) =========
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We thank the reviewer for bringing up several key points that will serve to improve the manuscript. We have addressed these concerns below.

General comments

Robinson et al. present a new ice sheet model. Using a zero-order hybrid SIA/SSA scheme, it is computationally inexpensive compared to higher-order and full-Stokes models. The manuscript is nicely written and provides a thorough description of the physics and its implementation in the model. In addition, the Yelmo model is available on a git repository with sufficient information to run it for a few standard configurations. The paper is worth publishing although I have a few comments and suggestions that could be considered.

Thank you for the very positive evaluation.

Main specific comments

- Most readers would probably like to see a more in-depth discussion on how the model perform for a real ice sheet configuration. While I applaud the authors for performing the EISMINT and MISMIP experiments since they are very informative on the model behaviour, I also think that the Antarctic ice sheet experiments are a bit weak and are briefly described (e.g. length of the simulation?). First, it could be nice to have more diagnostics in addition to the sole map of surface velocities (topography error but also other fields such as basal temperature / hydrology?). It is not necessary to show a perfect match with observations (which is most of the time achievable with a dedicated tuning), but it is interesting to see the bias structure to see if it is similar to other models with a similar complexity. Second, and more importantly, it could be very nice if you could discuss transient simulations. Since the model is suited for long integrations, some glacial-interglacial simulations (even with an idealised forcing) would be very interesting. If this is not possible, alternatively, you could maybe do the InitMIP experiments, since they are relatively easy to set up, and discuss your model results with respect to what is shown in Seroussi et al. (2019)?

We believe that more complex simulations fall outside the scope of this manuscript, as we would like to limit this model description paper to the ice-sheet model itself, and we expect more realistic simulations to be forthcoming soon in future studies. But we agree with the reviewer that more information about the Antarctica simulations could be provided. The present-day simulation is indeed comparable to the InitMIP experiment and the relevant metrics (rmse[H], rmse[vel], rmse[logvel]) already appear in the text. Nonetheless, we have added an additional plot explicitly showing the ice thickness bias and comparison of modeled velocity with observations, with relevant discussion.

- Unless I am mistaken, the model does not contain an isostasy model. This is certainly a limitation and might prevent its use for glacial-interglacial applications. Do you plan to account for this in the future?

Yelmo itself was designed specifically as a modular ice-sheet model (ice flow and thermodynamics). Consequently, it does not include an isostasy model, and we believe this is actually one of its important characteristics. Classically, ice-sheet models have been packaged as full system tools that contain several components performing isostatic, surface mass balance and shelf-melt calculations, among other things. Here we would rather like to publish an ice-sheet model in its minimal form. Numerous GIA models are available that can easily be coupled to Yelmo.

- Since you use an adaptive time step, a dedicated section could be very useful.

This is a good point. We have added a section to discuss the time stepping method and have added an additional figure for demonstration.

- Running Yelmo on my computer, I was not able to reproduce the results you show for the Antarctic ice sheet with the standard configuration file provided at the zenodo link. It is minor since I was simply digging for more info (basal drag coefficient value, trends, basal temperature etc.). Consider updating the configuration files for consistency with the results shown in the manuscript.

Thank you, this feedback is appreciated. The code archived at Zenodo for the published paper has been confirmed to work in all documented cases.

Other specific comments

- P11L22-25 Since beta=f(ub) for non-linear friction laws, does this mean that you have to do iterative loops (relaxation) to estimate beta for a given time step?

Yes, we iterate over beta and viscosity in a typical way. This is discussed in detail for beta when the SSA equations are introduced (P10L1).

- P14L13-20 The reader should be reminded that these experiments are SIA only.

Yes, this has been added.

- P15L8-10 Why is the motivation behind this choice of model parameters. You did the Antarctic experiments with a linear friction law, it would have been more useful to use the same model configuration for the two examples shown, wouldn't it?

The MISMIP benchmark experiments used these basal friction parameter values, so they are used here. In the Antarctica experiments, we wanted to keep the setup simple, so it was unnecessary to prescribe the particular values from MISMIP. We hope, as well, that it shows that it is easy to change from one law and parameter values to another.

Technical corrections

- P2L7 Typo (millenial).

Ok, this has been fixed.

- P2L7 Not only for palaeo, future multi-millenial change of ice sheets is also of interest

Yes, indeed, this has been changed.

- P7L9 n=3?

Correct, this has been fixed.

====== Short comment 2 (Ed Bueler) ==========================

We are grateful to Ed Bueler for providing critical comments regarding our work. We have considered all comments, and provide a point-by-point response below.

The paper under review is a model description of Yelmo, a new ice sheet model written in Fortran. The model is open source, and this reviewer successfully examined and downloaded the source from github, and compiled and ran the model, with unimportant technical difficulties only.

Yelmo has a very conservative design, with essentially no new physical features or submodels, at least as described here. If the publication standard at GMD is that the model is geophysical and is described by the submitted paper then this standard is unequivocably met. However, a basic description of a model, as part of its (evolving) documentation, should be part of its source code release. It is not clear that such model documentation is a publishable document; indeed a user's or developer's manual should match the changing versions of a code and address how specific capabilities are exploited.

It is true that the physics contained within Yelmo are rather conservative. However, the model design is rather unique, compared to many other models, and we believe this to be an added value to the community. A large section of the manuscript has been dedicated to explaining the model design from a technical point of view (not just numerics), which we believe is an important part of any model description paper and is often overlooked.

Because I suppose that the publication standard here is beyond that of such documentation, I have the following major concerns.

Concern 1. A new model should be justified by new directions for research, new ideas, and new capabilities. The most important concern: what is the

direction of this work? Substantial effort has been expended on Yelmo but it is not actually clear for what purpose. Open source ice sheet models exist with its capabilities, or with substantially greater capability (e.g. BiSCLES, CISM, ISSM, OGGM, PISM, Sicopolis), and all of these are forkable and to varying degrees modular. The authors of Yelmo chose to develop a new model, and not to add new capabilities to an existing model, despite some ideas coming from Sicopolis. So, where is this new one going?

One key phrase in this comment is "to varying degrees modular". It is not just modularity that is important (which is very important!), but also encapsulation. We built Yelmo explicitly to have a simple API that is intuitive and encapsulated (requires minimum definition of objects in external programs). Most notably, Yelmo should be quite capable for coupling with intermediate complexity climate models. We have tried to explain this motivation in the Introduction and Model design sections, but this has been further clarified in the revised manuscript. Nonetheless, we would point out that Yelmo has already proven extremely useful to us and we have noted interest from others, which indicates the demand for such a model, despite alternatives being available.

Concern 2. The "intended for collaborative development" and "flexible and user-friendly infrastructure ..." claims in the abstract, and repeated in various ways in the paper, are not demonstrated in any substantial way. For example, there is no demonstration that only minimal code extensions are needed to add a new capability. (Presumably this would be a great deal easier in e.g. Python than in Fortran anyway.) Does the model actually represent improved infrastructure for adding new capabilities? Noting that modularity does not, by itself, imply extensibility, if the model is extensible then the paper should demonstrate it.

This is a good point. However, it was difficult to find a simple example that wouldn't overly complicate the text. The proof of these statements, we believe, is nonetheless contained in the available source code.

Concern 3. The model verification mistakes of the past are recapitulated here. The EISMINT1 moving margin (MM) "benchmark" represented a failure of the community to read the literature 25 years ago, but there is no excuse now. As clear from textbooks (van der Veen, 2013, 2nd ed.) and well-known paper papers (Bueler et al, 2005), the Halfar (1983) exact solution is a full replacement for the MM experiment, which offers exact knowledge of what an SIA model should do. Regarding the EISMINT2 and MISMIP stuff, there is some excuse for using the benchmarks (though no evidence is given that the Yelmo runs offer more than the most common capabilities). The age- model testing via a (divide) analytic solution is applauded.

We agree that the EISMINT1/2 benchmarks are not true verification tests. However, they do serve as historical benchmarks and we believe can provide perspective on the performance of the model. Nonetheless, we have also implemented the Halfar/Bueler

exact test cases. Verification Test B in the Halfar configuration has been added in the revised manuscript.

Concern 4. Does the model run in parallel? The paper does not demonstrate or consider this but the tools seem to be suitable for it. In particular the Lis linear algebra library (ssisc.org/lis/), and the biconjugate gradient method, are used for the membrane stress solver component (SSA), and the library claims parallel performance. The concern here is that a lack of parallelism is exactly one of the limitations of many previous ice sheet models, thus limiting their attainable resolution.

The model has been built with parallelization in mind, however its capabilities have not been tested or fully implemented. The Lis library already supports parallel solving. The thermodynamics/enthalpy solver has been designed to be solved column-by-column, facilitating parallelization (as mentioned in the text). Additional smaller routines will also be easy to parallelize. Thus, this will not be a limitation of the model, but at this stage, it was not the first priority.

Now we turn from concerns to suggestions and questions.

The disadvantage of the GMD paradigm, of peer-reviewed publication of snapshot model descriptions, even users' manuals, is clear in this paper. Author and reviewer effort is devoted to a rapidly-out-of-date paper instead of (for example) devoting that effort to improving the software itself, or its evolving online documentation. Please see the mission statement of a different publication model: https://joss.theoj.org/about. The idea of the Journal of Open Source Software is to treat the software itself, and its evolving online infrastructure, as the reviewable object.

The raised discussion on publication paradigms is interesting and may be taken up at a different level. However, it seems rather outside the scope of this review to debate whether the GMD paradigm is better or worse than another journal. Having submitted to GMD, we do believe there is great value in a static and citable model description, which can serve as a common reference point for future work.

A surprise about the design of Yelmo was the decision to use only a temperature variable in the energy conservation equation. To a significant degree this means energy is not actually conserved. The alternatives, of course, include the replacement of temperature by an enthalpy variable, or the addition of a field to make a temperature/water-content pair; either makes possible energy conservation in polythermal ice. On the one hand there is no question that near-base polythermal ice is a concern for all ice sheet simulations; I would not want to see any serious treatment of ice sheet time scales without it. On the other hand, one would want the model to also be flexible enough to work for temperate mountain glaciers or Greenland outlet glaciers. The authors seem aware of this issue, but to have simply not

bothered. To the extent that there is a question here it is: what was gained by this choice?

Several models still choose to use temperature as the prognostic variable (CISM, GRISLI, IMAU-ICE, among others). Despite the lack of full energy conservation when water is present, this may not be a first-order issue for lower-resolution, long timescale simulations, given a number of other large uncertainties. In addition, the impact of water content on the rate factor is deeply uncertain. While an enthalpy solver would add a further degree of energy conservation, accurately mapping the transition from temperate to cold ice can only be achieved with very high vertical resolution, or potentially a two-layer scheme as in SICOPOLIS. In either case, this results in significant computational overhead and potentially numerical artifacts. We felt it was important now to have a fast, capable model that can run ensembles of long timescale simulations. In the revised manuscript we have added discussion of the energy scheme, and also show to what extent the temperature scheme can pass enthalpy benchmarks (Kleiner et al., 2015).

The above paragraph deliberately ignores the second-to-last sentence of the paper reporting a "plan to transition to" enthalpy, which only begs the question. Why not build the model based on the current state of the art from the beginning? This reviewer would be delighted to see a failed-or-not attempt to build a highly-principled and highly- conserving new model instead of a recapitulation of prior deficiencies.

See above. This is mainly a question of resources during the model development phase. As stated above, incorporating an enthalpy solver only brings added value when the transition from temperate to cold ice is well resolved. To do so in a computationally efficient manner requires careful consideration of the numerical treatment, which is reserved for future work. This is now discussed in the revised manuscript.

This reviewer had understood that Sicopolis (Greve) and GRISLI (Ritz) were models under active development which had capabilities roughly a superset of Yelmo. Is this true or not? Is this a fork of either model? (It seems not.) Is Yelmo already justified by its modularity and API design, somehow lacking in Sicopolis and/or GRISLI? It would help readers to expose these aspects of the design.

Both SICOPOLIS and GRISLI have active user bases and are undergoing development. The capabilities with respect to physics in all three models are rather comparable, but the model designs are quite different. We have extended the discussion of the model design choices made here that differentiate Yelmo from the others.

The Antarctica validation results are acceptable but suggest no capability which would draw-in researchers to use this model. To compare rather directly, a just-published paper suggests the power of many modern ice sheet models to account for the dynamics of the Antarctic ice sheet, namely the initMIP-Antarctica paper Seroussi et al 2019 (https://www.the-

cryosphere.net/13/1441/2019/). It lists 16 models of Antarctica (and 33 researchers), almost all of which would seem to have capabilities equal to or exceeding that of Yelmo. So where is this model going that is different, and why should we hope for new knowledge from it?

The goal of the Antarctica validation is simply to show that Yelmo can produce reasonable results for a realistic domain and for very different boundary conditions. Our results are in line with the models that appear in the initMIP-Antarctica study. Despite its similar physics to other models, it is sure that were Yelmo to be included in the ensemble, its results would not be identical. So there is potentially value in contributing to model diversity. However, Yelmo's main value at the current release is in its usability and transparency, as discussed above.

In summary, a new model like Yelmo, containing no significant new physics or model mechanisms, could in theory be useful. It could be a better piece of software than other offerings, it might have better performance, or it might be able to process input data faster or more flexibly, or it might just be implemented better and have better documentation. This paper does not convince me of any of it.

This conclusion is arguably subjective, and Yelmo will not be useful for everyone. However, we have made a concerted effort in the revised manuscript to make the advantages of Yelmo clearer.

Line-by-line, generally minor comments:

p 2 line 1: Lipscomb et al 2019 does not solve Stokes.

The phrasing here is ambiguous, and now has been corrected.

p 9 line 23: Hard-coding BCG for the SSA solve is a bad idea. Have there been experiments with AMG in Lis? Can a performance comparison be reported?

We have not performed additional sensitivity tests with variations on the Lis solver options. Through experience with SICOPOLIS, the BCG solver was recommended to balance speed and accuracy. We take the reviewer's point that the settings should not be hard coded, and the option has been added to the model parameters.

p 11 section 2.4: I am surprised by the temperature variable. If the claim is that converting to/from the enthalpy variable is too costly, then this should be stated.

We do not make that claim here. See above comments on enthalpy.

p 14 line 13: The EISMINT1 moving margin experiment has no justification *whatsoever*. Please use the Halfar (1983) solution so that you know the

exactly-correct prediction of the SIA. See "Test B" in Bueler et al (2005), and add "Test C" from that source if you want variety.

While we understand the reviewer's point, one justification we see of EISMINT1 experiments is that they serve as a historical reference. As mentioned above we have now added results from Test B to the revised manuscript.

p 14 line 16: "and thermodynamic": I hope not! EISMINT1 results depend on constant temperature (isothermal) ice, so energy will not be conserved.

The phrasing has been modified. While the feedback between temperature and dynamics was disabled in the EISMINT1 experiments, the temperature field was nonetheless diagnosed and can be used for comparison.

p 14 line 17: ""Type-I" discretization models": The fact that Yelmo agrees with other particular numerics is not relevant; EISMINT1 reported groups of results that way so as to expose a flaw not propose a standard. Instead, please take the opportunity to compare model results to exact predictions (a.k.a. analytical solutions) of the continuum model when available, *which they are in this case*. Beyond Halfar (1983) and Bueler et al (2005) for the SIA, there are exact solutions of the SSA, including results in van der Veen 1983), Schoof (2006), and Bueler (2014).

When comparing with EISMINT1 reported results, it is important to distinguish the numerical discretization, since as mentioned by the reviewer, there is no analytical solution to compare with. Again, we have added a comparison with Test B to compare with an analytical result.

p 14 line 29: This idea of smoothing is described in the reference Bueler et al (2007)as "non-physically 'smeared'". Whatever the meaning of this fiddle, it is not physics and the reference says that. (I think the point was that if *only* the temperature variable is symmetrized then the instability goes away, which modelers probably knew at the time but none had reported concretely. That is, the instability does not occur in a variable-softness model unless the softness variation is transported in 3D.) Supposing the EISMINT2 nonsense is valuable at all, please don't offer model users this smoothing, which hides physics.

We have removed this comment.

====== Reviewer comments 2 & 3 (Fuyuki Saito) =========

We thank Fuyuki Saito for the careful review and interesting code suggestions. Please find our response below.

This paper describes the numerical ice-sheet model Yelmo version 1.0. The Yelmo model is available on a git repository with sufficient information to run, and this manuscript contains mostly enough description of the model physics

and example application. I think this paper is fairly well written with some exception below, and can be accepted with minor revision.

(1) One point is about symmetry of the model (P14, L21 and after). Figure 3 is the simulated basal temperature of the experiments A and F of EISMINT2 configuration, and the paper states 'Yelmo produces symmetrical temperature patterns in both experiments,' First, minor one, I suggest to describe white kind of symmetry is the topic in this section. The configuration of EISMINT2 is 'radially' symmetric, but we all failed to simulate true radially symmetric pattern in particular for experiment F.

This is a good point. We are now more precise when discussing symmetry with respect to these tests.

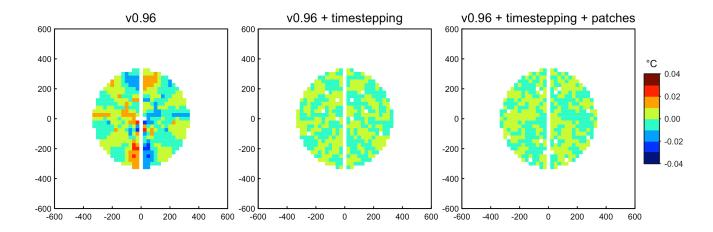
Second, major one: what is the degree of symmetry in the argument of this section? Actually, taking a closer look, some breaks of symmetries are already visible in the figure 3. The result of experiment A looks symmetric both along X and Y axis, but not along x=y diagonal. The result of F, even worse, shows breaks of symmetry both along X and Y axis (I attached a copy of figure 3 with marks to show the breaks of symmetry). So, even under the figure resolution, Yelmo already failed to produce symmetrical temperature patterns. In my opinion, preservation of model symmetries requires full control on the source and compilation, because even single change of arithmetic orders (e.g., (A+B)+C vs A+(B+C)) in a model can trigger and amplify breaks of symmetries. Yelmo depends on an external library in order to solve SSA equations, which is hardly controlled from outside, therefore Yelmo may find such symmetry breaking under an idealized configuration with ice shelves, even the SIA part is perfect. On the other hand, although preservation of the symmetries in the model is desired, it is not a top priority of a model, especially for one to simulate realistic worlds. We believe that such minor points have little influence on the simulation under realistic, highly asymmetric configuration for most of the application. So, I suggest the authors to keep the argument of Yelmo symmetries, and also state clearly the standpoint and/or main targets of Yelmo.

It is an important validation of the model to show that on a domain like that of EISMINT1/2, numerical artifacts do not arise leading to blatant asymmetry in the result. However, in any realistic cases, topographic and boundary forcing will likely dominate over minor asymmetries. So, it is indeed not a critical priority, given that Yelmo performs well in this regard. This has been clarified in the revised work.

It may not be a reviewer's work, I check the Yelmo source code to find the source of symmetry breaking. I attached some suggestion for Yelmo to preserve the numerical symmetries as a series of patch files since revision ed94c608516e2c46c7985ea98eea94fce47b37d8 (you can run git-am to apply them). It may be not complete and, honestly speaking, it may have bugs because I did not check in detail. The author can import if they like them, but

hope them to check the revision and results in detail before inclusion. If fortunate, SIA results will be more symmetric than before.

We kindly thank the reviewer for the extra effort of patching the code with these fixes. Independently, we also realized that the time stepping scheme with respect to the different components needed improvement for stability (moving from a simple Euler step to a predictor-corrector approach). This also resulted in significant improvement in the symmetry of the test. While applying your patches did improve the symmetry compared to the original model, in the updated version, the patches made no changes to an already satisfying result (see figure below, which shows the basal temperature field compared with itself mirrored along the y-axis; purely zero-error are white points).



(2) Another point is about precision (P4, L2). (See also minor points below for terminology of floating-point types).

The paper states that single and double precision give equivalent results, because the units of all time variables in Yelmo are cast in years instead of seconds and thus very small numbers are avoided. I do not understand this statement. I do not claim for the result but for the reasoning.

As far as all the quantities are larger than the smallest limit of floating-point number representation, same precision (significant digits) is kept either for the case with unit seconds and with unit years, because it differs not in precision but in the order of magnitude. The smallest number of a typical 'single' precision is around 1.18e–38. What variables do have possibility to show smaller value than this? The rate factor can be small, but even ice temperature is -100 degree Celsius, its magnitude is 1e-31 /s/Pa^3, which is large enough to be represented by single precision.

I agree that, a typical number of significant decimal digits of single precision is 7, which is actually smaller than the digits of the factor from year to second (31556926, 8 digits). If all the quantities in the model is originally defined with unit year, then it is possible to meet such situation, where unit-second version shows different results in the final digit. However, many parameters are originally defined in unit second and converted into unit year in the model,

thus round-off happens in some parameters themselves instead, which are almost the same situation as the unit-second case. (By the way, fortunately 31556926 can be fully represented by single floating-point number while 3155692[57] are not).

If there is a variable to be smaller than the limit in the case of unit-second, my question is solved. So, please give me an example.

Again but from a different point of view: this argument should depend on the model spatial resolution. As I mentioned, difference between single and double precision is merely the number of significant digits, if order of magnitude of all the quantities can be represented enough by the single. For a coarse resolution, difference in values at two adjacent grids of a field (e.g., surface elevation) is large enough to keep precision in their differences, However, for a higher resolution where the values at two become closer, so-called cancellation effects become large enough to reduce the precision of their difference. Relative error of the difference can be large enough for single precision to deviate from that by double precision. Generally speaking, a higher resolution experiment require high-precision computation to avoid such cancellation effects.

The concern here is primarily with avoiding extreme value divisions (large divided by small or small divided by large numbers), as well as so-called cancelation effects. This is particularly a concern in ice sheet modeling, where velocities, basal friction coefficients, effective pressure and rate factors can vary over several orders of magnitude. The statement in the manuscript was informed by experience. However, it is not critical to the manuscript, and so has been removed.

Minor points:

About precision (P4, L2) 'Single' and 'double' precision are in principle machine dependent characteristics although there are few exceptions. There is some definition of typical floating-point representation in IEEE754.

We note that in this sentence we equate single precision with 32 bits and double precision with 64 bits. This appears to be consistent with the definition in IEEE754, so we have left it as it is.

2.3, around Eq.(14). ub is defined as basal sliding above Eq.(14) while a depth-averaged velocity below (14). I am confused. Possibly typo?

In regions of plug flow that treated by the SSA solution, basal sliding and depthaveraged velocity should be identical.

2.4, below Eq. (27). Better to write as 'Horizontal diffusion is assumed negligible.'

Ok, this has been changed.

Table 1 last column. No degree mark.

Ok, this has been added.

There are two additional things to my first review.

First one, minor. There are contour lines in the figure 3, but what are they? I suppose they are surface elevation field. Anyway please explicitly describe them. I am sorry not to point out this in the first report. I just realized this now.

These are indeed contour lines. This will be added to the caption.

[Reviewer comment 3] Second one, also minor. I prepare two more patches in order to preserve symmetries. I confirmed symmetries of basal temperature field in X-axis Y-axis and diagonals at least for experiment F which was executed on my PC (compilation is by gfortran). Attached is a series of patches from revision ed94c608516e2c46c7985ea98eea94fce47b37d8, which includes all the patches I previously posted, so you can discard the previous one.

Thank you again. Please see the comments on symmetry above.

Description and validation of the ice-sheet model Yelmo (version 1.0)

Alexander Robinson^{1,2,3}, Jorge Alvarez-Solas^{1,2}, Marisa Montoya^{1,2}, Heiko Goelzer^{4,5}, Ralf Greve^{6,7}, and Catherine Ritz⁸

Correspondence: Alexander Robinson (robinson@ucm.es)

Abstract. We describe the physics and features of the ice-sheet model Yelmo, an open-source project intended for collaborative development. Yelmo is a thermomechanical model, solving for the coupled velocity and temperature solutions of an ice sheet simultaneously. The ice dynamics are currently treated via a "hybrid" approach combining the shallow-ice and shallow-shelf/shelfy-stream approximations, which makes Yelmo an apt choice for studying a wide variety of problems. Yelmo1000Yelmo's main innovations lie in its flexible and user-friendly infrastructure, which promotes portability and facilitates long-term development. In particular, all physics subroutines have been designed to be self-contained, so that they can be easily ported from Yelmo to other models, or easily replaced by improved or alternate methods in the future. Furthermore, hard-coded model choices are eschewed, replaced instead with convenient parameter options that allow the model to be adapted easily to different contexts. We show results for different ice-sheet benchmark tests, and we illustrate Yelmo's performance for the Antarctic ice sheet.

1 Introduction

The field of continental-scale, ice-sheet modeling started with a handful of pioneering models (e.g., Huybrechts et al., 1988; Ritz et al., 1997; Greve, 1997a). These models were computationally efficient for the resources available at the time. Typical grid resolutions were on the order of 20-40 km and generally the shallow ice approximation (SIA) was used to solve the ice dynamics. These classic models have been most useful for long time-scale paleo simulations in part because they are fast, but also because they are relatively simple in design, usually relying on low-tech solutions to numerical problems. Most of these models were designed before the era of the high-performance computing cluster, which made it challenging to build models otherwise.

Nowadays, a large number of ice-sheet models exist, supported by a growing and active community of developers. Models today represent a broad spectrum of approaches that incorporate different levels of physical complexity and computational ingenuity. These models include hybrid approaches that heuristically combine the SIA with the shallow shelf approximation (SSA)

¹Complutense University of Madrid, Madrid, Spain

²Geosciences Institute CSIC-UCM, Madrid, Spain

³Potsdam Institute for Climate Impact Research, Potsdam, Germany

⁴Institute for Marine and Atmospheric research Utrecht, Utrecht University, Utrecht, the Netherlands

⁵Laboratoire de Glaciologie, Université Libre de Bruxelles, Brussels, Belgium

⁶Institute of Low Temperature Science, Hokkaido University, Sapporo, Japan

⁷Arctic Research Center, Hokkaido University, Sapporo, Japan

⁸Institute for Geosciences and Environmental Research, Grenoble, France

(e.g., Bueler and Brown, 2009; Winkelmann et al., 2011; Goldberg, 2011; Pollard and DeConto, 2012; Pattyn, 2017; Quiquet et al., 2018) higher-order approximations (e.g., Goldberg, 2011; Cornford et al., 2013; Hoffman et al., 2018; Lipscomb et al., 2019), including full Stokes solutions (e.g., Larour et al., 2012; Gagliardini et al., 2013; Hoffman et al., 2018; Lipscomb et al., 2019)(e.g., Larour et al., Newer models often feature finite-element/finite-volume methods (e.g., Larour et al., 2012; Gagliardini et al., 2013; Hoffman et al., 2018) or adaptive mesh refinement (Cornford et al., 2013), which allows simulation of complex terrain and very high resolution where it is needed (e.g., at the grounding line in Antarctica). While more complex models are driving advances in our understanding of the physics and relevant processes of ice sheets over a range of time scales, simpler and thus faster methods are still required to understand the evolution of the ice sheets on multi-millennial, paleo-multi-millennial time scales.

Here we introduce the ice-sheet model Yelmo¹, which is intended to provide access to complex and robust model physics through an intuitive model design. It is a hybrid ice-dynamics model that is easy to use and configure. We expect that Yelmo will be useful for long time-scale paleo-simulations of the continental ice sheets, coupled climate – ice-sheet modeling, ensemble simulations and uncertainty studies, as well as for teaching. Below, we first describe the model structure, physics design (Section 2), followed by the physics (Section 3), timestepping approach (Section 4) and application programming interface (API, Section 25). Then, we present results for several benchmark experiments to validate the model performance (Section

2 Model design

(Section 58).

Yelmo has been inspired and largely derived from classical ice-sheet models that have been used successfully for many years — with the most in common with GRISLI (Ritz et al., 1997; Quiquet et al., 2018) and SICOPOLIS (Greve, 1997a, 2019). However, in contrast to many models, Yelmo was designed from scratch to run as a modular library that can be called by other programs rather than as a stand-alone executable. The strict application of this philosophy drove many design choices and allowed us to develop a robust ice-sheet model library with a clear API that would be difficult to develop in an ad-hoc way later. Thus, developing this framework was a primary reason to build a new model, rather than continuing the development of other active projects such as GRISLI and SICOPOLIS.

36), and a simulation simulation of the present-day and glacial Antarctic ice sheet (Section 47), followed by the conclusions

45Yelmo is written in Fortran90Fortran 2003, which provides continuity from previous code bases and supports the fact that clarity and readability of the code are important features. Like SICOPOLIS and other models, we have opted for "low tech" solutions whenever possible, meaning that internally coded routines are preferred and, thus, the external dependencies of the model are kept to a minimum. This ensures that the algorithms used remain accessible and easily changeable. Nonetheless, Yelmo has two key dependencies: the NetCDF library for convenient, community-standard input/output capability and the bobbrary of Iterative Solvers for linear systems² (Lis, Nishida, 2010), which is used for solving the elliptical SSA equations. The latter can be compiled with OpenMP parallelization active, which can speed up this computationally intensive step.

¹The name Yelmo refers to a semi-domed, rocky mountain in the Guadarrama Mountains outside of Madrid, Spain.

²https://www.ssisc.org/lis/

Yelmo has been designed to be user friendly (i.e., straightforward to understand), accessible, portable and adaptable. These features were facilitated by the design choice to separate what we call the "model accounting" from the model physics itself, and by following an object-oriented approach. There are no global variables in Yelmo (except for a few global constants related to the general physics of the planet being simulated), which means that variables and parameters are saved together in containers (called derived types in Fortran) specific to each of the main Yelmo components, such as dynamics or thermodynamics, as described in the sections below. These containers make up the individual components of the overall Yelmo object (itself a container) that contains all of the variables, parameters and information needed to simulate a given domain of the ice sheet. Multiple instances of the Yelmo object can therefore be defined in a program (e.g., one Yelmo-object instance for Greenland and one for Antarctica), and each one will operate fully in isolation from the others. This is the model accounting, which is of a specific design built into Yelmo and is thus the only part not easily portable to other models.

The model physics, meanwhile, consists of subroutines that are fully portable and, whenever possible, only rely on native data types (e.g., scalar, vector, arraysarray). In other words, the specific, non-portable design structure of the Yelmo object does not contaminate the physics subroutines, since the necessary variables and parameters are always passed as arguments. In general, This approach requires that all input and output to subroutines must be defined as arguments. Each argument must further always be given an intent characteristic (in Fortran, the intent of an argument can be one of IN, OUT or INOUT), which ensures that only the variables destined for output from the routine can be modified inside it. This approach not only aides debugging and provides programmatic safety, but provides a clear blueprint to users of what each subroutine does. Most importantly, the subroutines are thus fully self-contained and can be used in other programs and contexts, as long as the correct arguments can be provided.

Concerning the model accounting, the Yelmo object contains all parameters and variables needed to run a given domain. For clarity and convenience, it has been divided into four components: topography, dynamics, material properties and thermodynamics (Fig. 1). Each component has an associated set of functions to load parameters, allocate and initialize the variables, update the variables (i.e., the actual physics calculation step), and finally to terminate the instance of the component at the end of the program. This pattern is followed for all four components and represents the component-level API.

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Each component contains variables and parameters necessary for the calculation of its specific physics, however each component also relies on the variables defined in other components since the ice sheet is a highly-coupled and nonlinear system. The benefit of the somewhat artificial division of components made here is that the use of INTENT statements ensures that variables of a given component can only be modified in the corresponding module. For example, when the update subroutine of the topography module is called, only the object containing topography variables is defined as INTENT=INOUT, while the objects containing dynamics, material and thermodynamics variables are all defined as INTENT=IN. Analogous to the design of the physics subroutines, the use of intent statements here makes the model blueprint clear, but also enforces consistency with the overall design of the Yelmo structure. The hope is that this will not only make the model more user friendly, but it will also naturally lead to more disciplined model development in the future.

85In addition to the four components that contain prognostic and diagnostic model variables, the Yelmo object includes a boundary component, which defines all fields that Yelmo requires as input from external sources (Fig. 1). These fields can

be obtained from other coupled models, or simply by loading data, however Yelmo does not make any assumptions about their source. The boundary component is defined as INTENT=IN in all modules, so that Yelmo does not have the right to modify them internally. This conceptual isolation of the ice-sheet model serves to ensure that coupling with other models is as 90aightforward as possible.

, because it is clear by design which variables should be provided to Yelmo as boundary conditions. This is a key feature of Yelmo in comparison with many other models.

Yelmo also makes use of a working precision variable, which allows for the model to be compiled with any real precision. For most applications, single precision (32 bits) is sufficient. Double precision (64 bits) gives equivalent results for the tests we have made. This is in part due to the fact that the units of all time variables in Yelmo are cast in years instead of seconds and thus very small numbers are avoided. Nonetheless, this choice is left open to the user.

In terms of model physics, each component of Yelmo was built to work independently, in the sense that a given component is agnostic to the methods used to calculate variables from other components. For example, the temperature and velocity fields , which are taken as input to calculate the viscosity and rate factor in the material component, are used are used by the material component without any knowledge about the physics and numerical approximations used to calculate behind them. This means that sometimes simplifying assumptions cannot be used, even though they may be valid in some cases (such as assuming that the strain rate is only due to SIA terms where the ice sheet is frozen to the bed). However, the benefit is that typically the most general solutions possible have been implemented for each component. Thus, when the physics of one component is changed or upgraded, it is likely that the other modules will not require any modification.

Grid information is also stored in the main Yelmo object, and a single grid is defined for use with all components. Like many ice-sheet models, Yelmo uses the Arakawa-C grid staggering approach (Arakawa and Lamb, 1977) extended to 3D, as shown in Fig. 2. Scalar variables, such as temperature, are defined at the cell centers, which in Yelmo are designated as "aa-nodes". Velocity components and gradients are calculated on cell edges ("ac-nodes") and scalar coefficients, like diffusivity in the SIA approach, are calculated on cell corners ("ab-nodes"). The specific numerical discretization of the finite difference equations largely follows the approach of Macayeal (1997). The advantage of this approach is that it benefits from the natural staggering that occurs when calculating gradients (e.g., the surface slope is naturally defined on the ac-nodes), but it also results in greater numerical stability of the model (Macayeal, 1997).

Yelmo requires an evenly-spaced, Cartesian grid in the horizontal direction, while the vertical component follows a classic sigma-coordinate system (Greve and Blatter, 2009). The vertical axis ζ represents the relative height within the ice sheet, running from $\zeta = 0$ at the ice-sheet base to $\zeta = 1$ at the ice-sheet surface:

$$30 \quad \zeta(z) = H_z(z)/H \tag{1}$$

where z is the elevation relative to present-day sea level, $H_z(z)$ is the ice thickness up to the elevation z within the ice sheet and H is the total ice thickness. Yelmo can be defined with any specified number of vertical grid points, which can be unevenly spaced. Typically, we have set $n_z = 20$ and the ζ axis axis is defined with higher resolution near the base and surface of the ice sheet, which is important for resolving thermodynamics and ages accurately. Use of the sigma-coordinate system simplifies the

numerics of an evolving domain in the vertical direction and inherently results in higher resolution for grid points with less ice thickness (Greve and Blatter, 2009). Vertical velocities are calculated on ac-nodes in the vertical and aa-nodes in the horizontal, while horizontal velocities are calculated on ac-nodes in the horizontal and aa-nodes in the vertical. Boundary conditions in a vertical column are applied directly at the ice base and ice surface, which correspond to ac-nodes (see Fig. 2).

3 Model physics

Yelmo solves for two prognostic variables using coupled equations of mass and energy conservation: the ice thickness (2D field) and ice temperature (3D field). Velocity (3D vector field) is diagnosed from approximations of ice flow assuming a nonlinear flow law. These equations are described in the subsections below, along with additional considerations related to each component. For more details on the derivation of the equations, thorough explanations can be found in various references (Greve and Blatter, 2009; Cuffey and Paterson, 2010), and thus are not repeated here.

3.1 Topography

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The evolution of the ice thickness in the model is determined from mass conservation:

$$\frac{\partial H}{\partial t} = -\nabla \cdot H \bar{\mathbf{u}} + \dot{a} + \dot{b}_g + \dot{b}_f - \dot{c} \tag{2}$$

where H is the ice thickness, $\bar{\mathbf{u}} = (\bar{u}, \bar{v})$ is the depth-averaged horizontal velocity, \dot{a} is the surface mass balance, \dot{b}_g and \dot{b}_f are the basal mass balance for grounded and floating ice, respectively, and \dot{c} is the calving rate at floating ice margins. In Yelmo, in order to obtain more accurate mass balance accounting, the advection of ice and source contributions are treated separately as follows. First, a forward Euler explicit method (or optionally an upwind implicit method) is used to solve for the ice thickness at each timestep without accounting for \dot{a} , \dot{b}_g , \dot{b}_f or \dot{c} . The depth-averaged horizontal velocity is obtained from the dynamics component from the previous timestep. Note that dynamics is normally updated with the same timestep frequency as the topography component (see timestepping iteration (see Timestepping below). Next the mass balance terms \dot{a} , \dot{b}_g and \dot{b}_f are applied. It should be noted that the basal mass balance of floating ice is a boundary variable for Yelmo (i.e., it is obtained externally and passed to Yelmo), while the basal mass balance of the grounded ice is calculated internally as part of the thermodynamics solver (see Thermodynamics section below).

Yelmo also includes special treatment of grid points at the floating margin of the ice sheet, by making a distinction between ice-covered grid points that are totally and partially filled following Albrecht et al. (2011) and Lipscomb et al. (2019). This is done in a relatively simple, yet effective way to avoid artificially thin ice thickness at the ice margin. For each floating ice-covered grid point that has an ice-free neighbor, the reference ice thickness of the margin point (H_{ref}) is defined as the minimum thickness of the direct, ice-covered neighbors, or half of that in the case of grounded ice. This represents the minimum ice thickness for which the cell can be considered completely ice covered. The fraction of ice cover is then defined as $f_{ice} = \min(H/H_{ref}, 1)$. Whenever $f_{ice} < 1$, the grid cell is considered dynamically inactive and the ice thickness is stored in a buffer field during dynamic and thermodynamic calculations (i.e., these modules see the ice thickness as zero at these points).

Since the horizontal velocity components are calculated on ac-nodes, this method, which ensures zero ice flux through the downstream edge of a partially filled margin cell. During mass conservation, however, this buffer is added to the ice thickness field and the steps outlined above are applied. In this way, the ice cell can be filled with ice from upstream and when the threshold of $f_{\rm ice} = 1$ is reached, the ice sheet shelf can advance.

In the final mass conservation step, calving \dot{c} is treated at the floating ice margins. Currently, a simple threshold method has been implemented, as well as a threshold+flux method (Peyaud et al., 2007). In both methods, the calving rate applied to the ice sheet is defined following Lipscomb et al. (2019):

$$\dot{c} = \frac{H_{\text{ref}} - H}{\tau_c} \tag{3}$$

where τ_c is the characteristic calving time, usually set to 1-10 years, and $H_{\rm ref}$ is the margin ice thickness as defined above. Setting τ_c to higher values facilitates ice-shelf growth and thus grounding-line advance in transient, glacial simulations, but has little impact on the steady-state distribution of ice shelves for present day. This calving rate is applied only when the ice thickness of totally-filled-ice margin points is below a threshold value (simple threshold method), or when the ice thickness is below a threshold value and the upstream flux is not sufficient to return the ice thickness to above the threshold (threshold+flux method). For paleo simulations the latter is our preferred method, as it allows for more robust ice shelf advance (Peyaud et al., 2007).

As mentioned above, calving is only applied to floating margin points with $f_{ice} = 1$. However, a final check is also applied to both floating and grounded margin buffer points. If no neighbors are fully ice covered with $f_{ice} = 1$, then this buffer ice will also be calved and ice-margin retreat happens in a natural way.

Once the ice thickness has been completely updated, Yelmo diagnoses whether the ice should be grounded or floating. To facilitate this step, the distance from height above flotation as measured in ice thickness, i.e., how close a grid point is to the Archimedes flotation criterion, is calculated on each aa-node:

$$H_{\rm g} = H - \frac{\rho_{\rm sw}}{\rho} \left(\underbrace{\max}_{c} \left(z_{\rm sl} - z_{\rm b} \right), 0 \right) \tag{4}$$

where ρ is the ice density and $\rho_{\rm sw}$ the seawater density, and $z_{\rm sl}$ and $z_{\rm b}$ are the boundary fields of sea level and bedrock elevation, respectively. $H_{\rm g}$ can thus be positive, zero or negative. When $H_{\rm g}$ is positive, the ice thickness exceeds the flotation criterion, and is considered grounded, while when $H_{\rm g}$ is zero or negative, the ice is considered floating.

Yelmo also calculates the grounded fraction of each grid point, $f_{\rm g}$. On aa-nodes, $f_{\rm g}$ is only assigned binary values to maintain consistency with the overall grid definition: zero when $H_{\rm g} \leq 0$ or one when $H_{\rm g} > 0$. However, on <u>eell-edgesac-nodes</u>, the values of $f_{\rm g,acx}$ and $f_{\rm g,acy}$ are determined by linearly interpolating $H_{\rm g}$ from the two bounding aa-nodes. When both bounding aa-nodes are positive $f_{\rm g,ac} = 1$, and when both are negative $f_{\rm g,ac} = 0$. When one aa-node is positive $(H_{\rm g,pos}H_{\rm g,s})$ and one aa-node is negative $(H_{\rm g,neg}H_{\rm g,s})$, the grounded fraction on the ac-node is determined from linear interpolation:

$$(H_{g,\text{neg}g_{+}} - H_{g,\text{pos}g_{-}})$$

5 Alternatively, it is possible to calculate f_g via subgrid bilinear interpolation of H_g to intermediate points to determine the grounded area fraction. However, this operation is more computationally intensive, and we find that in practice, the simple linear interpolation method is sufficient.

The surface elevation (z_s) is calculated following Pattyn (2017) as

$$z_{\rm s} = \max \left[z_{\rm b} + H, z_{\rm sl} + \left(1 - \frac{\rho}{\rho_{\rm sw}}\right) H \right],\tag{6}$$

10 This approach ensures that the surface elevation solution is consistent with the Archimedes flotation criterion on aa-nodes.

The remaining tasks of the topography component are to diagnose other useful topographic characteristics, such as surface and ice thickness gradients (on ac-nodes) and topographic masks.

3.2 Material

The material component of Yelmo handles the calculation of the rate factor, the strain rate tensor and effective strain rate, the effective viscosity and, optionally, the age of the ice. Essentially, the material variables make the link between thermodynamics and dynamics, since the rate factor depends on temperature and the strain rate depends on velocity. No distinction is made between the type of approximation used to solve the dynamics here, rather all equations follow from the more general hydrostatic approximation (Greve and Blatter, 2009).

The effective viscosity, used to determine strain heating in the thermodynamics component, is calculated as

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$$\eta = \frac{1}{2} \left(\dot{\varepsilon}^2 \right)^{\frac{1-n}{2n}} \left(A^{-1/n} \right),$$
 (7)

where $\dot{\varepsilon}$ is the effective strain rate, n is the Glen's Flow law exponent (Glen, 1955; Greve and Blatter, 2009), typically set to n = 3, and A is the rate factor. The effective strain rate is given by the second invariant of the strain rate tensor ($\dot{\varepsilon}_{ij}$):

$$\dot{\varepsilon} = \left(\frac{1}{2}\dot{\varepsilon}_{ij}\dot{\varepsilon}_{ij}\right)^{\frac{1}{2}} \tag{8}$$

and the strain rate tensor itself, following index notation, is

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$$\dot{\varepsilon}_{ij} = \frac{1}{2} \left(\frac{\partial u_i}{\partial x_j} + \frac{\partial u_j}{\partial x_i} \right), \quad i, j = 1, 2, 3.$$
 (9)

The rate factor, A(x,y,z), can be prescribed to a constant value, or calculated as a function of ice temperature following an Arrhenius equation:

$$A(T') = E_f A_0 e^{-Q_a/RT'}$$
(10)

Here R is the ideal gas constant, A_0 and Q_a are the temperature-dependent rate factor coefficient and activation energy, respectively (see Greve and Blatter, 2009). E_f is a so-called enhancement factor, which is used to approximate the effect of anisotropic flow. In Yelmo, it is possible to specify different values of the enhancement factor for different flow regimes (shear, stream and shelf). The shelf value is prescribed anywhere ice is floating, while the inland value of E_f is a weighted average between the shear and stream value with the weighting given by a diagnosis of the vertical shearing fraction at any given point:

$$f_{\underline{\text{shear}}\mathbf{z}} = \frac{\left(\dot{\varepsilon}_{xz}^2 + \dot{\varepsilon}_{yz}^2\right)}{\dot{\varepsilon}^2}.$$
 (11)

Typical values of the enhancement factor for the shearing, streaming and shelf regime are $E_f = (3.0, 1.0, 0.7)$, $E_{\rm shr} = 3.0$.

0 $E_{\rm strm} = 1.0$ and $E_{\rm shlf} = 0.7$, respectively (Ma et al., 2010).

In addition, it is possible to track the deposition time (i.e., age) or other conservative tracers of the ice using the online an Eulerian tracer advection model. The general 3D advection equation of a conservative variable X,

$$\frac{\partial X}{\partial t} = -u \frac{\partial X}{\partial x} - v \frac{\partial X}{\partial y} - w \frac{\partial X}{\partial z},\tag{12}$$

is solved with a second-order, upwind explicit method. The ice surface boundary condition must be imposed. When tracing the ice deposition time, the ice surface boundary condition is X(t) = t. At the ice base, an initial deposition time is prescribed to be several thousand years before the start of the simulation, however this plays little role in the resulting vertical profile of deposition times. When ice is melting at the base $(\dot{b} < 0)$, the following flux boundary condition is defined (Rybak and Huybrechts, 2003):

$$\frac{\partial X}{\partial t} = -u_b \frac{\partial X}{\partial x} - v_b \frac{\partial X}{\partial y} - \dot{b} \frac{\partial X}{\partial z}.$$
(13)

Basal freeze-on is assumed to be negligible. It is well known that Eulerian solvers lose accuracy towards the base of the ice sheet, and therefore this method can only be considered to give a first-order estimate of age a conservative tracer (Greve et al., 2002; Rybak and Huybrechts, 2003). It can nonetheless be useful for diagnosing the age of ice, in order to know the timescale of different dynamic properties or to, e.g., impose an age-dependent enhancement factor (Greve, 1997b).

3.3 Dynamics

The Yelmo dynamics component is currently representative of a "hybrid" class of ice-sheet model, treating different modes of ice deformation via a combination of the simplifying shallow-ice and shallow-shelf approximations (SIA and SSA, respectively). In the following, the description of the dynamics equations follows closely the notation and definitions of Greve and Blatter (2009) and Pollard and DeConto (2012).

Yelmo treats the horizontal velocity u(x,y,z) and v(x,y,z) as the sum of transport via internal shear (u_i, v_i) and basal sliding (u_b, v_b) :

$$u = u_i + u_b$$

$$v = v_i + v_b.$$
(14)

Here, and analogously for v, $u_b(x,y)$ represents a depth-averaged velocity, so it is vertically constant, and $u_i(x,y,z_b) = 0$, where the subindex "b" here represents the basal boundary of the ice sheet. It also holds that in the vertical average (denoted by a bar), $\bar{u} = \bar{u}_i + u_b$. To calculate u_i and v_i , Yelmo uses zero-order SIA equations:

$$u_{i}(z) = -\left[2|\nabla \tau_{d}|^{(n-1)} \int_{z_{b}}^{z} A(z_{s} - z')^{n} dz'\right] \tau_{d,x}$$

$$v_{i}(z) = -\left[2|\nabla \tau_{d}|^{(n-1)} \int_{z_{b}}^{z} A(z_{s} - z')^{n} dz'\right] \tau_{d,y},$$
(15)

where $u_i(z)$ and $v_i(z)$ are the horizontal components of the SIA velocity as a function of depth at a given location, A is the material rate factor of the ice, which is obtained from the material component , and (Eq. 10), n is the Glen's Flow law exponent (Glen, 1955; Greve and Blatter, 2009) and $\tau_d = (\tau_{d,x}, \tau_{d,y}) = \rho g H \nabla z_s$ is the driving stress. In the horizontal plane, the term in brackets is calculated on the ab-nodes for stability and mass conservation (Type-I discretization) improved mass conservation (Huybrechts et al., 1996), and then it is staggered onto the ac-nodes where it is multiplied with the surface gradient driving stress. In the vertical plane, the horizontal velocities are calculated at the vertical center of each grid point (aa-nodes). Following Bueler and Brown (2009), we use the SSA solution to calculate the transport implied by sliding at the base (i.e., in regions of ice streams and floating ice shelves):

$$\frac{\partial}{\partial x} \left[\eta_d \left(4 \frac{\partial u_b}{\partial x} + 2 \frac{\partial v_b}{\partial y} \right) \right] + \frac{\partial}{\partial y} \left[\eta_d \left(\frac{\partial u_b}{\partial y} + \frac{\partial v_b}{\partial x} \right) \right] = \tau_{d,x} - \tau_{b,x}
\frac{\partial}{\partial y} \left[\eta_d \left(4 \frac{\partial v_b}{\partial y} + 2 \frac{\partial u_b}{\partial x} \right) \right] + \frac{\partial}{\partial x} \left[\eta_d \left(\frac{\partial u_b}{\partial y} + \frac{\partial v_b}{\partial x} \right) \right] = \tau_{d,y} - \tau_{b,y}.$$
(16)

where $(\tau_{b,x}, \tau_{b,y}) = -\beta(u_b, v_b)$ (or in vector notation $\tau_b = -\beta \mathbf{u}_b$) is the basal stress due to friction. The basal friction coefficient β is set to zero for floating ice shelves, and can otherwise be set to a constant value or follow another user-defined formulation (power law, regularized Coulomb, etc.), depending on the context . See the subsection on basal friction (see basal friction description below for details). The depth-integrated (2D) effective viscosity, which is only used for solving the SSA dynamics, is defined as

$$\eta_d = \left[\frac{1}{2} \left(\bar{A}^{-1/n} \right) \left(\dot{\varepsilon}_d^2 + \dot{\varepsilon}_0^2 \right)^{\frac{1-n}{2n}} \right] H \tag{17}$$

where \bar{A} is the vertically-averaged rate factor, $\dot{\varepsilon}_d$ is the 2D effective strain rate and $\dot{\varepsilon}_0^2$ is a small regularization factor for avoiding a potential singularity when velocity gradients are zero. The 2D effective strain rate is calculated as a reduced form of the second invariant of the strain rate tensor (Eq. 9) that does not include vertical shear terms:

$$\dot{\varepsilon}_d^2 = \left(\frac{\partial u_b}{\partial x}\right)^2 + \left(\frac{\partial v_b}{\partial y}\right)^2 + \frac{\partial u_b}{\partial x}\frac{\partial v_b}{\partial y} + \frac{1}{4}\left(\frac{\partial u_b}{\partial y} + \frac{\partial v_b}{\partial x}\right)^2. \tag{18}$$

In Yelmo, $\dot{\varepsilon}_d$ is only used for calculating η_d , while the 3D effective strain rate is calculated from the full strain rate tensor in the material component (see Material section above). Calculating the full tensor during the iterative SSA solution procedure would

be much more computationally expensive, while the 2D effective strain rate is already sufficient for the vertically integrated SSA equations (Pollard and DeConto, 2012).

The stress boundary condition imposed at the floating ice front, following Winkelmann et al. (2011) and Greve and Blatter (2009), is

$$\eta_{d} \left(4 \frac{\partial u}{\partial x} + 2 \frac{\partial v}{\partial y} \right) n_{x} + \eta_{d} \left(\frac{\partial u}{\partial y} + \frac{\partial v}{\partial x} \right) n_{y} = \left(\frac{1}{2} \rho g H^{2} - \frac{1}{2} \rho_{\text{sw}} g H_{o}^{2} \right) n_{x}
\eta_{d} \left(4 \frac{\partial v}{\partial y} + 2 \frac{\partial u}{\partial x} \right) n_{y} + \eta_{d} \left(\frac{\partial v}{\partial x} + \frac{\partial u}{\partial y} \right) n_{x} = \left(\frac{1}{2} \rho g H^{2} - \frac{1}{2} \rho_{\text{sw}} g H_{o}^{2} \right) n_{y}.$$
(19)

The depth of the seawater up to the flotation depth, H_o , is defined as: $H_o = \min\left(z_{sl} - z_b, \frac{\rho}{\rho_{sw}}H\right)H_o = \min\left(\max\left(z_{sl} - z_b, 0\right), \frac{\rho}{\rho_{sw}}H\right)$.

This is the depth of the ocean directly adjacent to the ice sheet, which acts to reduce the outward pressure at the floating ice margin. In constrast to Winkelmann et al. (2011), this boundary condition is not currently used in Yelmo for grounded ice, where Eq. 16 applies.

The SSA equations are nonlinear, elliptical, partial differential equations with non-local solutions. Yelmo uses Lis for the numerical solution using the biconjugate gradient method. The subroutine to discretize the equations and to call Lis was ported from the latest SICOPOLIS version 5-dev (Greve, 2019; Rückamp et al., 2019) and subsequently modified for model design choices in Yelmo. We use a Picard iteration method to account for the nonlinear dependence of the effective viscosity (η_d), and potentially the basal friction coefficient (β), on velocity. Convergence of the SSA solution is tested using the L^2 relative error norm (Gagliardini et al., 2013):

$$\delta_{\text{u,v}} = \frac{2\sqrt{\sum (u_1 - u_0)^2 + \sum (v_1 - v_0)^2}}{\sqrt{\sum (u_1 + u_0)^2 + \sum (v_1 + v_0)^2}},$$
(20)

where (u_1, v_1) and (u_0, v_0) are the velocity solutions for the current and previous iterations, respectively, and the sum is made over all grid points with non-zero velocity being considered by the SSA solver. By default, we consider a convergence limit of $\delta_{\rm u,v} = 10^{-2}$, which is typically achieved within 1-10 iterations, depending on the context. This limit can be specified by the user.

The result of solving the above equations is the hybrid, 3D horizontal velocity field (u, v). The vertical velocity w can then be diagnosed by applying a kinematic boundary condition at the base, and integrating the continuity equation for incompressible flow (Greve and Blatter, 2009), from z_b to z,

$$w(z) = \dot{b} - \left(u_b \frac{\partial z_b}{\partial x} + v_b \frac{\partial z_b}{\partial y}\right) - \int_{z_b}^{z} \left(\frac{\partial u}{\partial x} + \frac{\partial v}{\partial y}\right) dz'. \tag{21}$$

The vertical velocity is naturally defined on the ac-nodes in the vertical plane, analogous to the horizontal velocity in the horizontal plane. The above dynamics update results in a 3D hybrid velocity field (u, v, w) that is consistent with the current state of the topography.

3.3.1 Basal friction

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Basal frictional stress, as it appears in the SSA elliptical equations, is defined as

$$5 \quad \tau_b = -\beta \mathbf{u}_b \tag{22}$$

where β represents the basal friction coefficient, with units of [Pa yr m⁻¹], which can be defined in several ways. β is prescribed to be zero for floating ice, and otherwise can be set to a constant or a spatially varying field and, depending on the formulation used, it can depend on velocity itself. For this reason, we also define c_b as the bed friction coefficient, which we consider to only provide information about conditions at the physical bed (e.g., the nature of basal sediments, basal hydrology, effective pressure, etc.), independent of velocity. In the model, therefore, β is defined as:

$$\beta = c_b f(\mathbf{u}_b). \tag{23}$$

Thus in all formulations implemented in Yelmo, the term $f(\mathbf{u}_b)$ has units of [yr m⁻¹] and the coefficient c_b has units of [Pa], which helps to facilitate its physical interpretation.

Most commonly, β is defined using a linear (e.g. Quiquet et al., 2018), power-law (e.g. Pattyn, 2017), pseudo-plastic powerlaw (e.g. Aschwanden et al., 2013) or regularized-Coulomb (Joughin et al., 2019) formulation. The linear and power-law formulations are contained within the pseudo-plastic power-law formulation, so only the latter and the regularized-Coulomb formulation are needed to represent all four cases.

The pseudo-plastic power-law formulation (Schoof, 2010; Aschwanden et al., 2013) is

$$\tau_b = -c_b \left(\frac{|\mathbf{u}_b|}{u_0}\right)^q \frac{\mathbf{u}_b}{|\mathbf{u}_b|} \tag{24}$$

and thus $\beta = c_b u_0^{-q} |\mathbf{u}_b|^{q-1}$, with the pseudo-plastic exponent $q \in (0,1)$ and threshold speed u_0 . This expression results in purely plastic friction for q=0, linear friction for q=1 and power-law friction for 0 < q < 1. With q=1 and $u_0=1$, for example, $\beta = c_b$ and friction scales linearly with velocity. To obtain the power-law formulation used in the original MISMIP experiments (Pattyn et al., 2012), the following parameter values can be prescribed: q=1/3, $u_0=1\,\mathrm{m\,yr^{-1}}$ and $c_b=3.165176\times 10^4\,\mathrm{Pa}$.

Alternatively, the regularized Coulomb law (Schoof, 2005; Brondex et al., 2019; Joughin et al., 2019) is defined as

$$\tau_b = -c_b \left(\frac{|\mathbf{u}_b|}{|\mathbf{u}_b| + u_0} \right)^q \frac{\mathbf{u}_b}{|\mathbf{u}_b|} \tag{25}$$

and thus $\beta = c_b(|\mathbf{u}_b| + u_0)^{-q}|\mathbf{u}_b|^{q-1}$. Again q is the non-linear exponent and u_0 is an empirical threshold speed that dictates the transition from Coulomb friction when cavitation effects dominate at the base (typically for a hard bed) to Coulomb-plastic friction, when friction saturates (typically for weak till). When $u_0 = 0$ or q = 0, purely plastic friction is recovered.

The merits and physical basis of the different possible friction formulations and non-linear exponents are still part of under active debate (Aschwanden et al., 2013; Stearns and van der Veen, 2018; Brondex et al., 2019; Joughin et al., 2019), and all of the above formulations are used in ice-sheet modeling today. However, given the large uncertainty in boundary conditions

provided to an ice-sheet model, which include bedrock topography, sediment composition and distribution, basal hydrology and its temporal evolution, etc., it is clear that the use of any formulation will rely on empirical tuning. Also, as noted above, different choices for the friction exponents or threshold values can reduce a given formulation to another. Although modeling studies have shown that all four cases above can produce realistic velocity fields of the present-day ice sheets (e.g. Goelzer et al., 2018; Joughin et al., 2019), it remains to be seen how the choice of friction formulation may impact transient changes in the ice sheet.

For these reasons, we have chosen to implement the friction formulations in the most general way possible in the code, with essentially two free parameters: q as a non-linear exponent and u_0 as a threshold speed. Meanwhile, c_b is a 2D field that can backet to a constant value, or a spatially and/or temporally varying field based on e.g., whether the ice is frozen to the bed or temperate, on till strength (Bueler and van Pelt, 2015), effective pressure, or other user-defined criteria. As mentioned above, a Picard iteration method is used to solve for basal friction, η_d and the SSA velocity solution until convergence of the velocity solution is reached.

 β and c_b are initially defined on aa-nodes. c_b is naturally defined on the grid center, but while when $\beta = f(\mathbf{u}_b)$, the velocity components that are defined on ac-nodes must be staggered to the grid center. Once β has been calculated using one of the above formulations, it must be staggered to the ac-nodes for use in the elliptical solver. For purely floating points (i.e., $f_{\rm g} = 0$ at both bounding aa-nodes) $\beta_{\rm ac} = 0$, and for purely grounded points, $\beta_{\rm ac}$ is the average of the two neighbors. At the grounding line, Yelmo allows several options to handle staggering. These include simple averaging, taking the upstream value of β , taking the downstream value of β or taking the weighted average based on the grounded fraction of the ac-node.

3.4 Thermodynamics

Thermodynamics in Yelmo is treated in the classical way by solving the following energy conservation equation:

$$\frac{\partial T}{\partial t} = \frac{k}{\rho c} \frac{\partial^2 T}{\partial z^2} - u \frac{\partial T}{\partial x} - v \frac{\partial T}{\partial y} - w \frac{\partial T}{\partial z} + \frac{\Phi}{\rho c}$$
(26)

where k and c are the ice thermal conductivity and specific heat capacity, respectively. The <u>evolution of the</u> ice temperature T evolution is driven by vertical diffusion, horizontal and vertical advection, and internal strain heating due to ice shearing, Φ , where

$$\Phi = 4\eta \dot{\varepsilon}^2. \tag{27}$$

Horizontal diffusion is considered assumed to be negligible (Greve and Blatter, 2009). At the air-ice interface (i.e., the ice surface), the ice temperature is prescribed via the input boundary temperature field T_s , limited to a maximum value of $T_0 = 273.15K$. At the base of floating ice, the ice temperature is prescribed to the expected freezing temperature of seawater as a function of depth (Jenkins, 1991), except near the grounding line, where the temperature is prescribed to the pressure melting point of ice. At the base of grounded ice, when the ice temperature is below the pressure melting point, the vertical gradient of temperature is prescribed as $\partial T/\partial z = -Q_{\rm geo}/k$, where the geothermal heat flux ($Q_{\rm geo}$) is provided as a boundary field to Yelmo. If the temperature at the ice base reaches the pressure melting point, then the temperature is prescribed set to the

pressure melting point, and the basal mass balance is diagnosed as (Cuffey and Paterson, 2010):

$$5 \quad \dot{b}_{g} = -\frac{1}{\rho L} \left(Q_{b} + k \frac{\partial T}{\partial z} \Big|_{b} + Q_{geo} \right) \tag{28}$$

where $\dot{b}_{\rm g}$ is the basal mass balance of grounded ice (negative for melting), L is the latent heat of fusion for ice, Q_b is the basal heat production to due sliding and $\frac{\partial T}{\partial z}\Big|_b$ is the ice temperature gradient at the base. Yelmo provides calculates $\dot{b}_{\rm g}$ as which is a model output, and does not update the basal water layer thickness H_w internally, which can be achieved via external coupling with a basal hydrology modelin contrast to $\dot{b}_{\rm f}$ (basal mass balance of floating ice), which is prescribed in Yelmo as a boundary condition. Once the ice base is temperate (i.e., at the pressure melting point), it will remain so as long as $H_w = \begin{pmatrix} \rho_w & \dot{b}_{\rm g} \end{pmatrix} dt > 0 W_{\rm til} - \begin{pmatrix} \rho_w & \dot{b}_{\rm g} \end{pmatrix} \Delta t > 0$, where $\dot{b}_{\rm g}$ is used from the previous timestep and $W_{\rm til}$ is the water layer thickness in the till beneath the ice sheet. In other words, if it is expected that an energy deficit will result in freeze-on of the total available liquid water at the ice base, then the point is treated as a non-temperate ice point.

Yelmo simulates the evolution of the basal water layer thickness in the till following Bueler and van Pelt (2015):

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$$\frac{\partial W_{\text{til}}}{\partial t} = -\frac{\rho}{\rho_w} \dot{b}_g - C_d, \tag{29}$$

where C_d is the prescribed till drainage rate, usually set to $C_d = 0.001 \,\mathrm{m\,yr^{-1}}$. $W_{\rm til}$ is limited to the range $0 \le W_{\rm til} \le W_{\rm til,max}$ where maximum is usually set to $W_{\rm til,max} = 2 \,\mathrm{m}$. This approach allows for $W_{\rm til}$ to maintain consistency with the thermodynamic state of the ice sheet at all times. It does not include horizontal transport, as this could potentially be treated by an external basal hydrology model. It is also possible to disable calculation of $W_{\rm til}$ inside of Yelmo, and instead consider it as a boundary safetable. However, given the adaptive timestepping approach used by Yelmo, we have found that updating $W_{\rm til}$ internally at each timestep helps to avoid artificial oscillations that may develop otherwise when the thermodynamics and basal friction are coupled.

Eq. 26 is solved with an implicit method in the vertical direction, while the horizontal advection is solved separately applying an explicit, second-order upwind forward Euler method. This separation allows the energy conservation in the vertical to be solved as a 1D column model. The discretization of vertical diffusion follows the form presented by Hoffman et al. (2018), while the discretization of vertical advection follows a second-order central difference scheme. A given column of grid points consists of temperatures defined on the grid-centers (aa-nodes) and boundary values defined directly at the surface and base of the ice sheet.

3.5 Model interface

30 4 Timestepping

The Yelmo model-Yelmo makes use of a predictor-corrector (PC) method combined with adaptive timestepping to balance speed and stability, following the method developed by Cheng et al. (2017). This approach requires calculating the ice thickness twice per timestep, while all other variables can be calculated only once per timestep. Applying a PC method significantly

improves the accuracy of the solution compared to a simple forward Euler timestepping method. Furthermore, it facilitates the calculation of a stability metric at each timestep that can be used to evaluate model performance and forms the basis of a robust adaptive timestepping approach (Cheng et al., 2017). A given timestep therefore consists of three parts:

- 1. **Predictor step**: The topography component (namely the ice thickness) is predicted using the dynamics, material and thermodynamics solutions from previous timesteps.
- 2. **Update step**: Using the predicted topography solution, the dynamics, material and thermodynamics components are then also updated.
- 3. **Corrector step**: Using the updated dynamics, material and thermodynamics component solutions, the topography component is finally calculated again, starting from the ice thickness solution of the previous timestep.

In Yelmo, the predictor step is calculated via the second-order Adams-Bashforth (AB) method (Cheng et al., 2017),

$$H_{n+1}^{\star} = H_n + \underbrace{\Delta t_n}_{n} \left[\beta_1 f(H_n, \bar{\mathbf{u}}_n) + \beta_2 f(H_{n-1}, \bar{\mathbf{u}}_{n-1}) \right], \tag{30}$$

where H^* is the predicted ice thickness, Δt is the timestep, and $\beta_1 = 1 + \frac{\zeta_t}{2}$, $\beta_2 = -\frac{\zeta_t}{2}$ and $\zeta_t = \frac{\Delta t_n}{\Delta t_{n-1}}$. The labels n, n-1 and n+1 indicate the current, previous and next timestep, respectively. Here, $f(H, \bar{\mathbf{u}})$ is shorthand for $\frac{\partial H}{\partial t}$ as a function of the ice thickness and depth-averaged horizontal velocity field, noting that $\bar{\mathbf{u}}$ is also a function of the ice thickness, material and potentially thermodynamic state of the ice sheet. For this algorithm, β_1 , β_2 and ζ_t are timestep dependent, but the subscript n has been dropped for clarity. Once H^*_{n+1} has been calculated, the other components are updated, and finally the corrector step is then calculated via the Semi-implicit Adams-Moulton (SAM) method (Cheng et al., 2017).

$$H_{n+1} = H_n + \frac{\Delta t_n}{2} \left[f\left(H_{n+1}^{\star}, \bar{\mathbf{u}}_{n+1}\right) + f\left(H_n, \bar{\mathbf{u}}_n\right) \right], \tag{31}$$

where H_{n+1} is the corrected ice thickness for the next timestep.

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For the AB-SAM timestepping method, Cheng et al. (2017) have derived the following expression for the leading term of the local truncation error:

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$$\tau_{n+1} = \frac{\zeta_t \left(H_{n+1} - H_{n+1}^* \right)}{(3\zeta_t + 3)\Delta t_n}$$
 (32)

The local truncation error is valuable for diagnosing the performance of the model, and can be used as an indicator of numerical stability. For a small enough timestep, H_{n+1}^* and H_{n+1} should be indistinguishable and $\tau^{n+1} \sim 0$. However, as the timestep increases, the local truncation error will also increase.

An adaptive timestepping approach based on a proportional-integral (PI) controller method is therefore used to maximize the timestep while maintaining the truncation error below a specified threshold (Cheng et al., 2017; Söderlind and Wang, 2006). Defining the maximum truncation error over all grounded grid points as $\eta = \max |\tau|$, the next timestep is calculated using the

so-called PI4.2 controller Söderlind (2002):

$$\Delta t_{n+1} = \left(\frac{\epsilon}{\eta^{n+1}}\right)^{(k_I + k_p)} \left(\frac{\epsilon}{\eta^n}\right)^{-k_p} \Delta t_n,\tag{33}$$

where ϵ is the target tolerance and $k_I=2/10$ and $k_p=1/10$ are reasonable control parameters for the second-order AB-SAM 415 estepping method used here (Söderlind and Wang, 2006). This algorithm ensures that the time step increases when $\eta < \epsilon$ and decreases when $\eta > \epsilon$. The use of both η^{n+1} and η^n helps to avoid rapid fluctuations in the timestep, which improves model stability and results in a predictable timestep size as a function of the target tolerance.

For practical purposes, the timestep is further treated as follows. The timestep must be larger than a user-prescribed minimum value, but smaller than the Courant–Friedrichs–Lewy (CFL) 2D advective limit:

$$\Delta t_{\rm cfl} = C_{\rm cfl} \, \max \left| \frac{\bar{u}}{\Delta x} + \frac{\bar{v}}{\Delta y} \right|^{-1} \tag{34}$$

where the maximum is taken over all grid points and $C_{\rm cfl}=1.0$. Furthermore, the adaptive timestep is adjusted to ensure that the model stays synchronized with the frequency that Yelmo is being called externally. We found that the latter requirement often results in highly uneven timestepping; e.g., if Yelmo is called with a timestep of $\Delta t_{\rm tot}=2.0\,{\rm yr}$ and the first adaptive timestep is determined to be $\Delta t_1=1.9\,{\rm yr}$, then the second timestep would likely be $\Delta t_2=0.1\,{\rm yr}$. To avoid this possibility and increase substility, the condition is imposed that if any given adaptive timestep is predicted in the range of $0.5\Delta t_{\rm tot}<\Delta t<\Delta t_{\rm tot}$, then $\Delta t=0.5\Delta t_{\rm tot}$. In this example, this condition would ensure that $\Delta t_1=\Delta t_2=1.0\,{\rm yr}$, unless Δt_2 needed to be smaller for stability. Finally, if the maximum local truncation error η is larger than a specified threshold for any given integration, then the integration is discarded and repeated with a progressively smaller timestep until η diminishes and stability is restored, or the timestep reaches the minimum allowed value.

5 Model interface

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The Yelmo model interface is designed to be clear and simple, but also flexible. In its essence, there are three main model functions: yelmo_init to initialize the model variables, yelmo_update to perform the ice-sheet model calculations for a given timestep and yelmo_end to terminate the Yelmo object (free it from memory).

The first subroutine, yelmo_init, is used to load parameters, initialize variables in memory (i.e., allocate arrays) and, optionally, to initialize the topographic state variables (ice thickness, masks, etc.). No other variables are initialized here in the sense of being populated with data values, which is left to the user. An additional, optional helper function can be used, yelmo_init_state, which populates the remaining model variables in the material, thermodynamics and dynamics components. This initialization step is separated from that of topography because in practice, sometimes boundary variables (e.g., surface temperature) need the surface elevation as input in order to be determined. In contrast, the remaining variables, namely dynamics and thermodynamics, often rely on boundary variables to be initialized. Thus, a typical initialization sequence for a stand-alone ice-sheet model simulation could first call yelmo_init, then load or calculate boundary variables and then

call yelmo_init_state to finalize the initialization of all Yelmo variables. After this sequence, the Yelmo state should be consistent with running the model for one timestep with the prescribed boundary conditions and a fixed topography. If the model will be initialized from a restart file, then these data are loaded in each case based on parameter choices.

The next subroutine, yelmo_update, is used to advance the model state to a new timestep. Any modifications to boundary variables are left to the user externally, and Yelmo expects that the boundary conditions are valid for this timestep. The subroutine does not take any arguments to modify the model behavior – rather, all model configuration choices are specified in the parameters of the Yelmo object itself. These are initially loaded from a parameter file in the call to yelmo_init, however, it is possible to modify any parameter values during simulations, allowing for changing model configuration the model configuration transiently depending on the experimental setup. An additional optional subroutine, yelmo_update_equil, is available to facilitate equilibration. This routine effectively calls yelmo_update for a specified time window with unchanging boundary conditions, and allowing for the temporary modification of some key model parameters (such as the maximum allowed adaptive time step and the maximum allowed SSA velocity).

The last subroutine, yelmo_end, simply removes the Yelmo object from memory (i.e., all domain variables are deallocated). After calling yelmo_end, it is possible to reinitialize the Yelmo object via yelmo_init, for example, in order to test a different grid resolution or other configuration.

There are several input/output routines defined for Yelmo. yelmo_write_init can be used to initialize a NetCDF model output file with the axes of model dimensions defined from the Yelmo object and writing of static fields like domain masks. The writing of model output for inidividual timesteps is left to the user to maintain flexibility, as most programs require specific fields to be written (examples can be found in the test programs included with the code - see further below). In addition, yelmo_restart_write will create a NetCDF file and write all Yelmo fields as a snapshot, which can be used to restart the model (loading of a restart file can be activated with parameter choices).

As mentioned above, given the object-oriented approach, it is possible to run multiple Yelmo domains in one program. Each domain must be initialized separately via yelmo_init, and the variable fields populated with initial values, then separate calls to yelmo_update are needed during timestepping, and finally each object should be terminated at the end of the program via yelmo_end. With this structure, minimum modification of another model, like a global climate model is needed, to incorporate online ice-sheet evolution, or to simulate an ensemble of ice sheets in one program. Furthermore, all fields are directly accessible within the main program to facilitate coupling. For example, the 2D array of surface elevation of the topography component of the Yelmo Antarctica domain could be referenced as yelmo_ant%tpo%now%z_srf. While it is clear that the nesting of several containers (derived types) results in a rather long variable reference, it is unambiguous and straightforward to use.

6 Model validation and benchmarks

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Yelmo has been tested against several ice-sheet model validation tests and benchmarks in wide use today. These include the Halfar dome experiment (Halfar, 1983; Bueler et al., 2005), the EISMINT1 (Huybrechts et al., 1996) and EISMINT2 (Payne

et al., 2000) model intercomparison experiments that test uncoupled and coupled dynamics-thermodynamics, respectively, and MISMIP (Pattyn et al., 2012) for ice-shelf dynamics, among others. By design, many of these experiments allow isolation of specific model features for testing. When the model passes more complex benchmark tests, the simpler experiments are somewhat redundant (if the model passes a coupled thermodynamics-dynamics benchmark, the model should necessarily also be able to pass a dynamics-only benchmark). However, it should be noted that in the process of model development, all tests prove to be extremely useful. The results of all tests will not be reported here, but several are highlighted below to demonstrate that Yelmo performs well.

The Halfar dome experiment, a specific case of the more general Test B of Bueler et al. (2005), tests the ice-sheet model dynamics using the SIA solver alone. This test consists of simulating a radially-symmetric ice-sheet with zero mass balance and resting on a flat bed, deforming under gravitational stress. The analytical solution is known at every time, allowing a direct 485nparison of the simulation to the desired result. The simulation parameters consist of the margin radius and dome elevation, in this case set to the values suggested by Halfar (1983): $R_0 = 21.2132 \,\mathrm{km}$ and $H_0 = 707.1 \,\mathrm{m}$ – see Bueler et al. (2005) for further details. Figure 3 shows the root mean square error (RMSE) of the simulation with the analytical result after 200 years for a range of model resolutions. Yelmo demonstrates first-order (p = 1.01) numerical convergence with resolution towards the analytical result.

The EISMINT1 moving margin experiment also tests the ice-sheet model dynamics using the SIA solver alone, with an imposed constant rate factor and diagnosed thermodynamics (i.e., thermodynamics do not impact the ice-sheet configuration). Radial steady-state surface mass balance and background surface temperature fields are imposed as boundary conditions. Starting from ice-free conditions, the ice sheet simulated by Yelmo grows to dynamic and thermodynamic equilibrium within 25 kyr and 100 kyr, respectively. The steady-state summit elevation of Yelmo is 2992.43006.6 m compared to the reported range of 2997.5±7.4 m for so-called "Type-I" discretization models like Yelmo (where diffusivity is staggered to the ab-nodes). The basal temperature relative to the pressure melting point (i.e., homologous temperature) at the summit simulated by Yelmo is -12.85-13.37 °C, which lies within the EISMINT1 range of -13.40±0.56 °C. Other These and other relevant statistics are given in Table 1.

We also use the EISMINT1 moving margin experiment to demonstrate the capability of the adaptive timestepping approach 500Yelmo. By setting the tolerance parameter ϵ , Yelmo automatically adjusts the timestep to maintain the truncation error in ice thickness η around this value. Figure 4a shows the time series of the adaptive timestep used by Yelmo for a 25 kyr simulation for different resolutions. The timestep exhibits oscillations around a mean value, which is typical for such a PID approach (Cheng et al., 2017). When the timestep grows larger, the truncation error increases. This leads to a reduction in the timestep and the error decreases. Figure 4b shows the mean timestep used by Yelmo over the last 10 kyr of the simulation 505sus model resolution. Given a tolerance of $\epsilon = 10^{-2}$, Yelmo's mean timestep is $\Delta t = 6.96$ yr, $\Delta t = 1.59$ yr, $\Delta t = 0.24$ yr and $\Delta t = 0.06$ yr for resolutions of 50 km, 25 km, 10 km and 5 km, respectively. As expected, the timestep must be reduced for higher resolutions. These results are in line with those of Cheng et al. (2017) for the same experiment ($\Delta t = 12.4$ yr for 60 km resolution). It should be noted that the truncation error increases non-linearly as a function of the timestep, so setting a higher tolerance does not translate directly into a larger timestep.

510Next we validate the thermodynamics component first by performing the benchmark experiments Test A and Test B of Kleiner et al. (2015). In contrast to an enthalpy solver, Yelmo uses a temperature solver that assumes all water produced in the ice column drains directly to the bed and so temperate ice in the vertical column has no water content. In cases where water content of up to 3% could be present in the basal layers of the column, Yelmo's solver would be inaccurate. Nonetheless, we expect that the temperature solver should be sufficiently accurate to simulate ice sheets on long timescales and large spatial domains. Figure 5 shows the performance of Yelmo's temperature solver for Test A of Kleiner et al. (2015), which simulates a column of ice in a parallel-sided slab with no horizontal advection and no internal strain heating that undergoes warming and enthalpy solver should give identical, energy-conserving results. Yelmo's basal melt rate is essentially identical to the analytical solution for this problem and its transient behaviour is robust.

520n contrast, Fig. 6 shows the results of Yelmo's temperature solver for Test B, which simulates a parallel-sided slab on a sloping bed with a prescribed horizontal velocity and strain heating profile in steady state. In this case, water is generated in the basal layers of the ice column (see Fig. 6c), however Yelmo cannot reproduce this solution. Nonetheless, because temperature is limited to the pressure-melting point, the simulated ice temperature profile is in full agreement with the analytical profile. This is true both for a very high resolution case ($\Delta z = 0.5 \,\text{m}$) and a lower resolution case ($\Delta z = 10 \,\text{m}$), which allows us to 505 clude that the performance of the temperature solver is robust.

The EISMINT2 benchmark experiments A and F are useful for testing the thermodynamically coupled ice-sheet model with SIA dynamics like in EISMINT1. The experiments are identical to the EISMINT1 moving margin experiment, except the resolution is doubled (25 km) and the surface temperature is prescribed to be independent of ice thickness. Experiment A prescribes a summit temperature of 238.15 K, while experiment F is 1515.00 K colder, which promotes an increase in the region of ice frozen to the bedrock. The statistics for these experiments are listed in Table 1 as well. Figure 3-7 shows the basal homologous temperature distribution for experiments A and F. Yelmo produces symmetrical temperature patterns in both experiments, which are consistent with both the benchmark results (Payne et al., 2000) and other more recent models (e.g., Bueler et al., 2007; Hoffman et al., 2018). Yelmo also Axial symmetry, assessed by comparing the basal temperature field with a mirror of itself along the x- or y-axis, is maintained to a precision of 10⁻² K. This symmetry is not critical to realistic applications, but a lack of at least axial symmetry in this test is often indicative of numerical artifacts. In experiment F. Yelmo produces the so-called "cold spokes", which have been shown to be related to internal strain heating in regions of steep gradients in ice thickness, and largely numerical in nature (Bueler et al., 2007).

— Yelmo includes an option for smoothing the strain heating field as suggested by Bueler et al. (2007), however its effectiveness has not been tested yet in detail.

We also test the capability of the SSA solver and grounding-line treatment by running the MISMIP protocol experiments (Pattyn et al., 2012). Particularly, MISMIP EXP 1 (advance) and EXP 2 (retreat) are useful for testing the reversibility of grounding line advance, given the bedrock is defined as a linear downward sloping bed. The rate factor is prescribed according to steps that first decrease, allowing grounding line advance, then increase back to the original value. According to theory (Weertman, 1974; Schoof, 2007), only one steady-state grounding line position should exist for each step 1000- i.e., the ice

sheet should advance and retreat symmetrically without showing hysteresis. It is now well known, however, that ice-sheet models at coarse resolutions (1 km and greater) are unable to capture proper grounding-line migration, even when subgrid parameterizations to mimic higher resolution are applied (Seroussi et al., 2014; Gladstone et al., 2017).

In the MISMIP experiment performed here, the linear, downward sloping bedrock is defined in the x-direction as $z_b = 720 - 778.5 (x/750.0)$ with x in km and z_b in m. The bedrock elevation does not change in the y-direction, which extends to ± 50 km to allow the simulation of a symmetric ice stream flowing in the positive x-direction. The power-law formulation of Eq. 24 is used with the parameter values q = 1/3, $u_0 = 1$ m yr⁻¹ and $c_b = 3.165176 \times 10^4$ Pa. The rate factor is initially prescribed to $A = 1 \times 10^{-16}$ Pa⁻³ yr⁻¹ and the simulation is run for 25 kyr to equilibrate. Next, the rate factor is stepped evenly in log-space every 10 kyr until reaching $A = 1 \times 10^{-19}$ Pa⁻³ yr⁻¹, and then the rate factor is increased in the same way until returning to the original value.

Figure 4.8 shows results for this MISMIP experiment with Yelmo at different grid resolutions, ranging from 20 km down to 2.5 km, and with different treatments of basal friction near the grounding line. When the default model setup is used, with no special treatment at the grounding line, the grounding line advance is consistent for all resolutions. However, none of the lowest resolution simulations show grounding line retreat as the rate factor increases again. At a resolution of 5.05 km, some minor grounding line retreat can be seen, and for 2.5 km, the model is more successful at retreating though it remains 400 km from the target. In contrast, when the basal friction β is scaled at the grounding line by the grounded fraction of the ac-node ($f_{g,ac}$), the hysteresis is greatly reduced. The 5 km simulation retreats to within 200 km of the original position and the 2.5 km simulation retreats to within 100 km of the original position, thus showing convergence to the correct solution with resolution. With this setup, even the 10 km and 20 km simulations retreat significantly. In a third case, the basal friction is also linearly scaled to zero as the ice sheet approaches flotation (Leguy et al., 2014; Gladstone et al., 2017). In this case, the hysteresis and differences between different resolution simulations are further reduced, however, the system also tends to advance much less given all other conditions are the same.

Yelmo's Eulerian ice-age conservative tracer model is validated with a simulation of ice age in an idealized cofiguration against the analytical solution at the summit presented by Rybak and Huybrechts (2003). In this case, summit-like conditions are imposed, in that horizontal advection is neglected, and the vertical velocity is assumed to decrease linearly with depth. Fig.Figure 5-9 shows the solution with Yelmo as compared to the analytical result. For a nominal vertical resolution of $n_z = 30$ points and single or double precision, the age tracer gives errors of around in the range of 0.2% over the entire depth 0.5% over most of the column of the ice sheet, with higher errors at the base. Increasing the vertical resolution to $n_z = 50$ points decreases the error by an order of magnitudeand reducing the vertical resolution to $n_z = 10$ points increases the error to about 1%, while using $n_z = 30$ with higher resolution at the base of the ice sheet allows a similar reduction in error with significant computational savings. For Eemian-age ice in such simplified conditions, this the latter case gives an uncertainty of about-less than 1 kyr. It is expected that the error would increase for more realistic domains (Rybak and Huybrechts, 2003)3D domains, however the Eulerian age solver can be used for a first-order estimate of the age-depth profile in the ice sheet (Rybak and Huybrechts, 2003).

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7 Antarctica

880 further validation of the model's performance, we ran steady-state simulations of the present-day and glacial Antarctic ice sheet. These simulations, run at 32 km resolution, have been deliberately simplified to include the minimum complexity necessary to simulate the ice sheet without additional external components. There was no active isostasy model and geothermal heat flux was set to 50 mW m⁻² everywhere. The bedrock topography and initial ice thickness were prescribed from the RTOPO2.1 dataset (Schaffer et al., 2016). Basal friction followed a linear law, where $\beta = c_f \lambda_b (\rho g H) \beta = (c_f \lambda_b / u_0) (\rho g H)$ such $u_0 = 100 \, \text{m yr}^{-1}$ used as a scaling constant. We prescribed $c_f = 2 \times 10^{-3} \, c_f = 0.15$ (unitless) for most of the domain, except for ad-hoc adjustments in specific regions to improve the match with observations. This was additionally scaled by an exponential function of bedrock elevation: $\lambda_b = \min[1.0, \exp((z_b - z_1)/(z_1 - z_0))]$, where $z_0 = -200$ analogous to the approach of (Martin et al., 2011). We set $z_1 = 250 \, \text{m}$ and $z_1 = 0$ everywhere and $z_0 = -2000 \, \text{m}$, which ensures that the basal friction decays to low values for marine-based sectors of the ice sheet (Martin et al., 2011) for WAIS regions feeding the Ronne 590 shelf and $z_1 = -200 \, \text{m}$ elsewhere. Friction was scaled by the grounded fraction at the grounding line, but no additional scaling is applied. The enhancement factor parameters set for these simulations were $E_{\text{shr}} = 2.5 \, E_{\text{stron}} = 0.7 \, \text{and} \, E_{\text{shif}} = 0.5$. The bedrock topography and initial ice thickness were prescribed from the RTOPO2.1 dataset (Schaffer et al., 2016), after which the model ran for 50 kyr, reaching a steady-state modeled ice distribution.

For the simulation of the present-day state, surface mass balance (SMB) and surface temperature boundary fields were pre-**505** bed from a RACMO2.3 simulation driven by ERA-INTERIM data and averaged over 1981-2010 (van Wessem et al., 2018). The ice-shelf basal mass balance was set to a spatially constant value of -0.1-0.2 m a⁻¹. Figure 7 shows—where floating ice exists today and to -2.0 m a⁻¹ elsewhere. Figures 10 and 11 show a comparison of the simulations with the observed topography (RTOPO2.1) and the present-day observed velocity (Rignot et al., 2011). With this relatively simple model setup, it is nonetheless-possible to obtain reasonable agreement with observations. The root mean square errors (RMSEs) in ice thick-66%, velocity and log(velocity) are $\frac{290}{320}$ and $\frac{230}{320}$ m yr⁻¹ and $\frac{1.8}{1.9}$ log[m yr⁻¹], respectively, which fall in the range of other models in the initMIP-Antarctica intercomparison project (Seroussi et al., 2019). The dome configuration, slow deformational speeds and even most ice streams as they penetrate inland are well represented by the model, simulated ice sheet is thinner than the observed ice sheet over large parts of East Antarctica, with a broad positive bias near the South Pole (Fig. 11). The margins of the ice sheet are the most difficult to match and, in particular, the grounding-line positions of the 60% ice shelves. This is likely due in large part to the use of a spatially constant value for ice-shelf basal mass balance, but also reflects the simple approach used to represent β . Nonetheless, the results are, leading to larger biases in these regions. This pattern is quite consistent with other models (e.g., Martin et al., 2011; Quiquet et al., 2018; Seroussi et al., 2019). studies (e.g., Martin et al., 2011; Quiquet et al., 2018; Albrecht et al., 2020). Overall, the dome configuration, slow deformational speeds and even most ice streams as they penetrate inland are well represented by the model (Figs. 10 and 11).

We use the same setup with modified boundary conditions to simulate a configuration resembling that of a deep glacial period like the Last Glacial Maximum. The surface temperature was set to 10° C colder and the present-day SMB was maintained, except for points with a low or negative SMB were prescribed with a minimum value of $0.1 \, \text{m a}^{-1}$. The ice shelf basal mass

balance was set to a spatially constant value of $0.00 \,\mathrm{m}\,\mathrm{a}^{-1}$ and sea level was lowered by 120 m. In this case, the grounded ice sheet advances until the continental shelf break and thickens inland (Fig. 710). A similar structure of ice streams can be seen, due to the topographic dependence of β , but their speed is greatly reduced compared to those of the present-day simulation. We do not expect this configuration to be realistic, given that isostasy plays no role and a present-day-like SMB has been imposed. However, this test demonstrates that Yelmo is capable of resolving continental-scale changes in the ice sheet configuration in a plausible way.

8 Conclusions and future work

We have described the features and physics of the hybrid ice-sheet—shelf model Yelmo. Yelmo includes the physics to simulate continental-scale ice sheets and floating ice shelves using "shallow" approximations of the ice dynamics. The fully coupled thermomechanical ice-sheet model has been validated against several benchmark tests, and has been shown to simulate the dynamic configuration of the Antarctic ice sheet well.

Yelmo is expected to be useful for long-time scale simulations and/or ensembles. It is particularly suited for easy coupling with other models. For example, the simulation of multiple ice-sheet domains with independent parameter configurations coupled to a global climate model can be achieved in a simple and straightforward way. Also, given that the subroutines representing the physics of the model have been isolated from the "model accounting", it is possible to test individual model components in different contexts easily. This should facilitate future model development and comparison of different methods.

The model framework has been designed to facilitate the incorporation of new and different physics. Thus, this initial release of Yelmo lays the foundation for several future developments. These may include more advanced calving and basal friction schemes, as well as improved treatment of the grounding line. We also plan to transition to an enthalpy-based thermodynamics solver, and however this will require an adaptive vertical axis to be able to map the height of transition between temperate and cold ice accurately. We also plan to implement a variationally-derived "depth-integrated-viscosity approximation" solver (following e.g., Goldberg, 2011; Pollard and DeConto, 2012; Lipscomb et al., 2019) in the near future.

10 Code availability. Yelmo is maintained as a git repository hosted at https://github.com/palma-ice/yelmo under the licence GPL-3.0. Model documentation can be found at https://palma-ice.github.io/yelmo-docs/. The exact version of the model, along with the necessary input data, used to produce the results used in this paper is archived on Zenodo (https://www.doi.org/10.5281/zenodo.3782650) and has been tagged in the repository as v1.02.

Author contributions. A.R., J.A.S. and M.M. conceived the model design and features. A.R. wrote the model code with contributions from the remaining authors. All authors contributed to the model testing and writing the manuscript.

Competing interests. Heiko Goelzer is a member of the editorial board of the journal.

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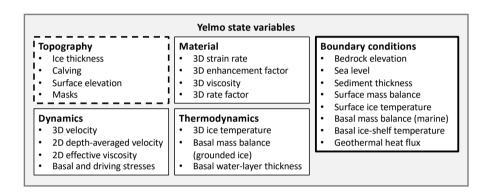


Figure 1. Overview of the Yelmo model structure highlighting state variables in the four components: topography, dynamics, material and thermodynamics, as well as the boundary conditions required to run the model. The thick black border for boundary variables indicates that these fields are never modified internally by Yelmo, while the components with a thin black border or dashed line are allowed to be modified depending on the context. When, for example, the topography is updated (dashed line), no other components are allowed to be modified (solid lines).

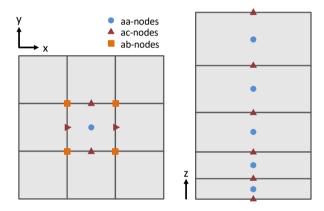


Figure 2. Yelmo staggered grid definition and nomenclature. The horizontal grid (left) assumes constant resolution in the x- and y-directions, while in the vertical (right) variable resolution is allowed. With any given cell defined as a 3D box, scalar variables are calculated on cell centers (aa-nodes), velocities are calculated on cell faces (ac-nodes, edges in 2D), and scalar coefficients are calculated on cell edges (ab-nodes, corners in 2D). Figure design adapted from Hoffman et al. (2018).

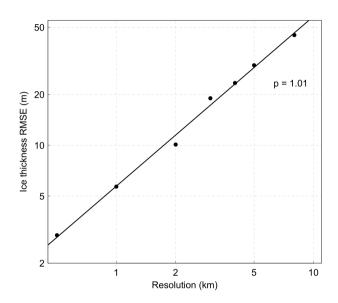


Figure 3. Steady-state, basal homologous temperature Root mean square error ($^{\circ}$ CRMSE) distribution of the Halfar dome experiment after 100200 kyr obtained yr simulated by Yelmo in EISMINT2 test A (left) and test F (right) compared to the analytical solution versus model resolution. Areas that have reached The value of p=1.01 indicates the pressure-melting point have been shaded greyorder of convergence as the resolution increases.

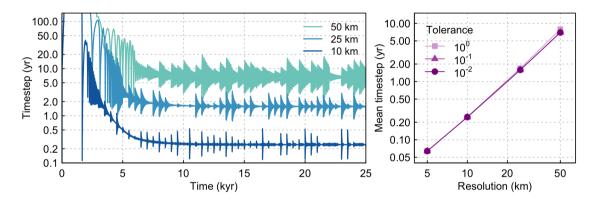


Figure 4. Adaptive timestepping for the EISMINT1 moving margin experiment. Time series of the timestep used by Yelmo for grid resolutions of 50 km, 25 km and 10 km and a tolerance of $\epsilon = 10^{-2}$ (left), and the mean adaptive timestep in the time range of 15-25 kyr versus model resolution (right). Separate lines in the right panel show results for different values of the tolerance parameter ϵ .

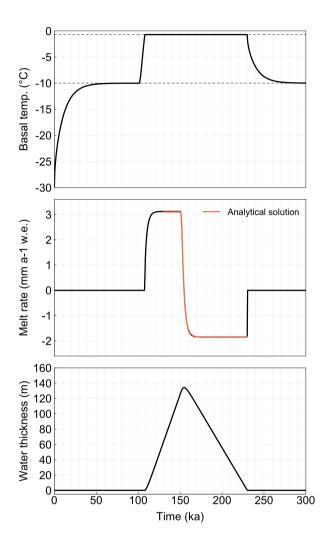


Figure 5. Time series of the basal temperature (top), basal melt rate (middle) and basal water layer (bottom) corresponding to the thermodynamic benchmark experiment Exp. A (Kleiner et al., 2015). The analytical solution (thick, light-red line) for the basal melt rate is compared to Yelmo results (black lines). Not that where the Yelmo results are not visible, they overlap with the analytical solution.

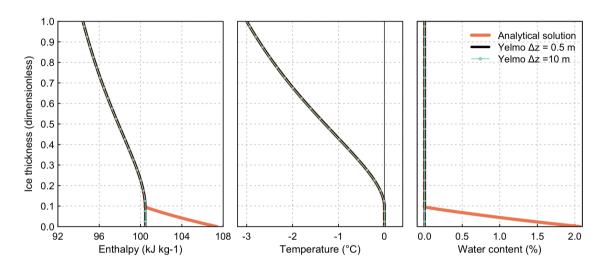


Figure 6. Steady-state vertical profiles of enthalpy (left), temperature (middle) and water content (right) for the thermodynamic benchmark experiment Exp. B (Kleiner et al., 2015). The analytical solution (thick, light-red lines) is compared to Yelmo results for a vertical resolution of $\Delta z = 0.5$ m ($n_z = 400$, black lines) and $\Delta z = 10$ m ($n_z = 20$, light green lines). The vertical, grey line in the middle panel shows the pressure-melting point as prescribed in this experiment. Note that where the analytical solution is not visible, it overlaps with the Yelmo results.

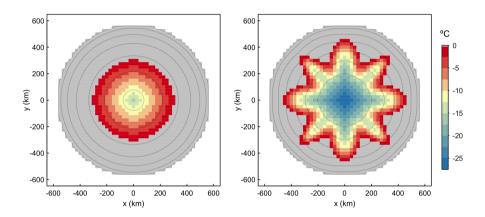


Figure 7. Steady-state, basal homologous temperature (° C) distribution after 100 kyr obtained by Yelmo in EISMINT2 test A (left) and test F (right). Areas that have reached the pressure-melting point have been shaded grey. The contour lines represent ice thickness at 500 m intervals up to 3500 m.

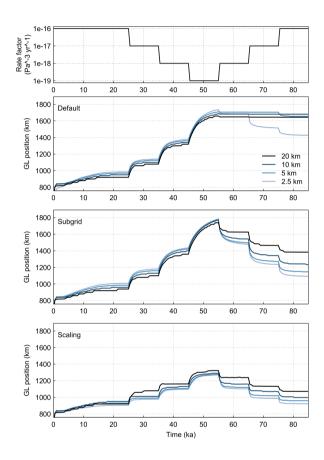


Figure 8. Yelmo performance in the MISMIP bedrock advance and retreat simulations on a linear sloping bed. The top panel shows the imposed rate factor A, with 10 kyr steps of decreasing and then increasing values. The three lower panels show the grounding line position evolution for each of three model configurations, respectively: "Default" is the standard model setup, with no special treatment of friction at or near the grounding line, "Subgrid" uses the grounded fraction at the grounding line to scale the basal friction, "Scaling" applies both the grounded fraction, and imposes a linear reduction in basal friction as the ice sheet approaches flotation. Separate simulations were run for resolutions ranging from 20 km down to 2.5 km.

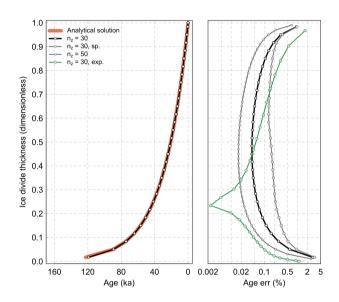


Figure 9. Age-depth Analytical age-depth profile for idealized summit compared to Yelmo Eulerian age tracing. The ice age relative to present day (left) with an is shown for the analytical solution (thickgrey, light-red line) and Yelmo (black line) with a vertical resolution of $n_z = 30$ with a linear vertical axis and compiled at double precision, and the associated error (right, %black line). The associated relative error (right) is given for additional resolutions this case (black line), as well as for a higher resolution of $n_z = 10$ and $n_z = 50$ are also shown and for comparison $n_z = 30$ compiled at single precision (thin dark grey lines, smaller points), and finally for the same a resolution of $n_z = 30$ compiled at double precision, but single precision with exponentially increasing resolution at the base instead of a linear axis (thin, light grey green line, smaller points).

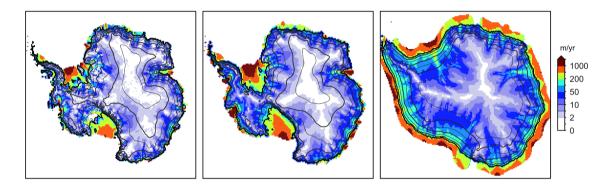


Figure 10. Antarctica present-day ice sheet configuration and surface velocities from observations (left), compared to a steady-state simulation with Yelmo (middle). In contrast, Antarctica glacial configuration and surface velocities simulated by Yelmo (right). Simulations were performed at 32 km resolution. The colors show surface velocity in m yr $^{-1}$ and the dark grey contours show surface elevation in 500 m intervals (thick lines correspond to $\frac{0}{2}$, 1000, 2000 and 3000 m above sea level). The black line shows the grounding line position.

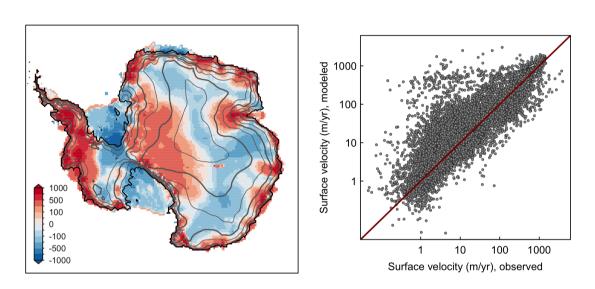


Figure 11. Simulated present-day ice-thickness minus observations (left) and simulated versus observed ice surface velocity (right). The colors in the left panel show the ice thickness difference in m and the dark grey contours show surface elevation in 500 m intervals (thick lines correspond to 1000, 2000 and 3000 m above sea level). The black line shows the grounding line position. In the right panel, the dark red line indicates a perfect correlation between model and observations.

Table 1. Yelmo performance in the EISMINT1 moving margin experiment ("Moving"), as well as in the EISMINT2 experiments A and F. Where available, metrics with the ensemble mean and standard deviation from the original benchmark experiments are also provided for comparison.

Experiment	Model	Volume [10 ⁶ km ³]	Area [10 ⁶ km ²]	Melt fraction	Divide thickness [m]	Divide basal temperature [K]	Divide homologous basal temperature [°C]
Moving	EISMINT1 Yelmo	-	- 1.003	- 0.58 <u>0.66</u>	2997.5 ± 7.4	_	-13.40 ± 0.56
	Tenno	1.939-1.980			2992.6 3006.6	256.9 -257.2	-13.61 -13.37
A	EISMINT2	2.128 ± 0.073	1.034 ± 0.043	0.72 ± 0.15	3688 ± 48	255.6 ± 1.4	-
	Yelmo	2.205 <u>2.170</u>	1.031	0.71 - <u>0.75</u>	3736 - <u>3678</u>	254.9 -254.7	-15.03-15.26
F	EISMINT2	_	_	_	_	_	_
	Yelmo	2.458 <u>2.373</u>	1.040_1.031	0.36 0.55	4374-4266	240.9 -240.7	-28.50 -28.80