

Using `trullekrul` in `pde2path` – anisotropic mesh adaptation for some Allen–Cahn models in 2D and 3D

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Abstract

We describe by means of some examples how some functionality of the mesh adaptation package `trullekrul` can be used in `pde2path`.

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1 Introduction

The Matlab package `pde2path` [UWR14, Uec19c] is designed for numerical continuation and bifurcation analysis of systems of PDEs of the form

$$M\partial_t u = \nabla \cdot (c \otimes \nabla u) - au + b \otimes \nabla u + f, \tag{1}$$

where $u = u(x, t) \in \mathbb{R}^N$, $t \geq 0$, $x \in \Omega$, $\lambda \in \mathbb{R}^p$ is a parameter vector, $M \in \mathbb{R}^{N \times N}$ is a mass matrix, which may be singular, and the coefficients c, a, b and f may depend on x, λ and u . Details on the terms in (1), the discretization of (1) by the FEM, the boundary conditions associated to (1), and how to use `pde2path` to compute branches of steady and time periodic solutions of (1), can be found in [RU19, Uec19b, Uec19a] and some further tutorials which together with the software and demos can be downloaded at [Uec19c].

Here we explain how to use the anisotropic mesh adaptation package `trullekrul` [JG16, Jen17] in `pde2path`. For this we extend the introductory tutorial [RU19] by a number of some advanced examples for steady Allen–Cahn type problems, i.e., problems of type

$$G(u) := -c\Delta u + f(u), \quad u = u(x) \in \mathbb{R}, \quad x \in \Omega \subset \mathbb{R}^d \text{ a bounded domain}, \tag{2}$$

with diffusion constant c , 'nonlinearity' (everything except diffusion) $f : \mathbb{R} \rightarrow \mathbb{R}$ (which also depends on parameters). For Ω we shall restrict to rectangles (2D, $d = 2$) and cuboids (3D, $d = 3$), and f and the boundary conditions will be given for the specific examples below.

In [RU19] we also explain some settings for mesh adaptation for problems of type (2) based on mesh refinement using standard a posteriori error estimators, in 1D and 2D. In the context of solution branches $s \mapsto (u(s), \lambda(s))$ for (2) as computed by the main user–interface routine `cont` of `pde2path`, where λ is used as a symbol for the active continuation parameter, it is desirable to adapt the mesh during continuation, for instance after a number of continuation steps, or if the error estimate is above some user given threshold. This mesh *adaptation* so far has been done in `pde2path` in a simple ad hoc way: first we *coarsen* the current mesh to a given (essentially fixed and uniform) background mesh by interpolation of the current solution to the background mesh, and then generate a new mesh by *refining* the coarse mesh and solution. The coarsening is necessary because otherwise we may end up with (at places) unnecessarily fine meshes. However, the simple approach sketched above is not very efficient, may lead to undesired branch–switching after coarsening, and, moreover, has so far not been fully implemented in 3D. By interfacing `pde2path` with `trullekrul`, we now have genuine 2D and 3D *adaptation* options during continuation of branches, which means coarsening (only) where appropriate together with moving of mesh–points and refinement.

In §2 we explain the general setup of `trullekrul` in `pde2path`, based on the standard data structures of `pde2path` with all data stored in the struct `p`, i.e., function handles to the rhs of (8a) and its Jacobian, FEM mesh, file-names for saving, controls for plotting, numerical constants such as stepsizes and solution tolerances), and the solution `p.u` itself and the current tangent to the solution branch. For this background, the `pde2path` data structures, the continuation algorithms, and the general usage of `pde2path`, we refer to [dWDR⁺19] and the tutorials at [Uec19c], in particular [RU19]. In §3 we then explain the usage of `trullekrul` by some example problems of type (8a).

2 General setup of `trullekrul` in `pde2path`

Given a function $u : \Omega \rightarrow \mathbb{R}$, `trullekrul` aims to optimize the FEM mesh to minimize the interpolation error $\|u - u_h\|_{L^p(\Omega)}$, where u_h as usual is the continuous piecewise linear interpolation of the nodal values. This is based on the discrete Hessian $H(u_h)$ of u_h , and the associated metric field

$$\Psi = \frac{1}{\eta} (\det(\tilde{H}))^{-\frac{1}{2p+d}} \tilde{H}, \quad (3)$$

where \tilde{H} denotes a matrix of absolute values of the eigenvalues of $H(u_h)$, and η is a scaling factor, which can be used to control the number of mesh points. In [CSX07] it is shown that meshes that are uniform wrt to Ψ minimize $\|u - u_h\|_{L^p(\Omega)}$. Thus, using Ψ , edge lengths L_{elem} of the triangulation are computed wrt to the metric Ψ , and then elements are coarsened/refined according to the following algorithm (cf. [LSB05, Fig.6]):

The details of each of the steps 1 to 3 in Algorithm 2.1 are controlled by a number of `trullekrul` parameters, which in `pde2path` we store in the field `p.trop` (`trullekrul options`) of the basic data struct `p`. To initialize `trop` we provide the two functions `troptions2D` and `troptions3D`, which as variants of the `trullekrul` function `gen_options` set `trullekrul` parameters in the way which appear to be most robust and efficient in 2D, respectively 3D. For us, the most important parameter is η in (3), where, for fixed L_{up} , larger η gives less triangles to refine. For adaptation during continuation we typically want to keep the number of mesh points n_p below some bound, and thus η should be set by some function which depends on n_p . To give maximum flexibility, the user may provide such a function in `trop.etafu` with signature `eta=etafu(p,np)`. The default setting is `trop.etafu=@stanetafu`, which returns the constant $\eta = 10^{-3}$, while a simple dependence on n_p is given by `eta=etafua(p,np)` which (by default) returns $10^{-5}np$. Altogether, in Table 1 we list the `trop` parameters/function handles which we find most useful for tuning the adaptation, and we strongly recommend to experiment with these and the other `trullekrul` parameters, see the sources for detailed comments. The `pde2path` parameter to

Algorithm 2.1: Outline of `trullekrul` mesh adaptation. Given a current solution u_h , the inner adaptation loop is performed, and the resulting new u_h is used as an initial guess for a Newton loop on the new mesh. This outer step is repeated `p.nc.ngen` times.

Compute the maximum edge length L_{\max} in metric space, and perform the following loop until $L_{\max} < L_{\text{up}}$ or until `it=imax`:

1. Eliminate the edges shorter than L_{low} by coarsening the respective elements; this includes swapping of elements with wrong orientation.
2. Refine the elements with $L_{\text{elem}} > L_{\text{up}}$ by splitting their longest edge and postprocessing (splitting of adjacent elements);
3. Move mesh-points according to a smoothing algorithm based on the (discrete) Laplacian.
4. Update Ψ, \tilde{H} and hence L_{elem} and L_{\max} .

switch on mesh-adaptation by `trullekrul` instead of option (i) within `cont` is `p.sw.trsw > 0`.

In some examples (in particular in 3D, see below), it is useful for mesh adaptation within `cont` to first use only the coarsening and moving functionality of `trullekrul`, and then adapt again with refinement. For this two step approach we provide a modification of the main `trullekrul` wrapper function `adapt_mesh`, named `tradapt`. Both, `adapt_mesh` and `tradapt`, take the option field `trop` as last argument, and for `tradapt` this should contain the field `trop.sw` which encodes the operations

$$\text{(face or edge) swapping, coarsening, moving (of mesh points), and refinement,} \quad (4)$$

in a binary way according to Table 2. Thus, coarsening and moving as a preparatory step¹ before adaptation is encoded as `p.trcop.sw=5`, and additionally `p.trcop.npb` should contain the desired maximum number of meshpoints.

3 Example implementations and results

To illustrate the use and performance of the `trullekrul` mesh adaptation we discuss some demos from `pdepath/acsuite`, which also collects the demos discussed in [RU19], and again we stress that new users should at least briefly browse [RU19, §4] and the associated demos.

3.1 2D

Extending ac2D from [RU19]. We start with (2) on the rectangle $\Omega = (-2\pi, 2\pi) \times (-\pi, \pi)$, with Dirichlet BC, i.e.

$$G(u) := -c\Delta u - \lambda u - u^3 + \gamma u^5 = 0 \text{ in } \Omega, \quad (5a)$$

$$u = d \cos(y/2) \text{ on } \Gamma_2 := \{x = 2\pi\}, \text{ parameter } d, \text{ and } u = 0 \text{ on } \partial\Omega \setminus \Gamma_2. \quad (5b)$$

The label $\Gamma_2 = \{x = 2\pi\}$ is due to the `pde2path` convention that the boundaries of rectangles as generated by `stanpdeo2D` have the order bottom-right-top-left. For $d = 0$, (8) features the bifurcation

¹executed if `p.trcop.npb > 0` and `p.trcop.cmax > 0` by calling `tradapt(..., p.trcop)`, i.e., with the 'trullekrul coarsening options' `p.trcop` instead of the 'trullekrul options' `p.trop`

Table 1: Most important parameters/function handles in `p.trop`, see also `troptions2D` and `troptions3D` for further parameters and comments. `p.trcop` (see below) needs the same parameters as `p.trop`, and additionally `p.trcop.sw`, see bottom of table. Typically, we just copy `trop` to `trcop`, set `p.trcop.npb`, and reset selected parameters such as `p.trcop.sw`.

Parameter	meaning, default value
<code>etafu</code>	default <code>eta=stanetafu(p,np)</code> returning 0.001. Larger <code>eta</code> gives less elements to adapt. See also <code>etafua</code> , yielding $\eta = 10^{-5}n_p$, and we recommend to copy <code>etafua</code> to the local directory and experiment with the prefactor.
<code>zfu</code>	function handle with signature <code>z=zfu(p)</code> , to select the field u for which the interpolation error is estimated. Default <code>zfu=@stanzfu</code> for which $z = u_1$ (first component of current solution). May be useful to overload for multi-component systems. In some cases, also scaling of z is useful, e.g., $z = \exp(u_1)$.
<code>setids</code>	function handle needed to link the <code>pde2path</code> data structures with <code>trullekrul</code> in case that different boundary segment numbers (and different BCs on different segments) are assigned in <code>pde2path</code> . Defaults: <code>@setidssq</code> in 2D (corresponding to a rectangular domain, as also used in <code>stanpdeo2D</code>); <code>@setidsbar</code> in 3D (corresponding to a cuboid domain, as also used in <code>stanpdeo3D</code>).
<code>sw</code>	behavior of <code>tradapt</code> according to Table 2; default 15.
<code>ppar</code>	p to optimize the interpol. error in the p -norm metric. Default: 1000, i.e., close to ∞ norm.
<code>innerit</code>	number of iterations in <code>trullekrul</code> , default 2. (Not to confuse with <code>p.nc.ngen</code> giving the 'outer' number of adaptation iterations).
<code>Llow, Lup</code>	lower/upper thresholds for edges in metric space, defaults $1/\sqrt{2}$ and $\sqrt{2}$. Smaller <code>Llow</code> can be used to avoid too much coarsening.
<code>qualP</code>	weight for combining mesh quality in metric space and euclidean space. Defaults: 0 in 2D (metric space only), 2 in 3D (avoiding too acute tetrahedra)
<code>trcop.npb</code>	desired number of mesh-points for pure coarsening steps.
<code>trcop.cmax</code>	maximum number of pure coarsening steps.

points

$$\lambda_{jl} = (j/4)^2 + (l/2)^2, \quad \phi_{jl} = \sin(j(x + 2\pi)/4) \sin(l(y + \pi)/2), \quad j, l = 1, 2, \dots \quad (6)$$

from the trivial branch $u \equiv 0$. The associated bifurcating branches have already been discussed in [RU19, §4.1] and are computed in `ac2D/cmds1`. In `ac2D/cmds2` we redo some of these computations starting with a very coarse mesh and aiming to illustrate the use and performance of `trullekrul`. Some results are shown in Fig. 1, which compares the older mesh adaptation by error estimators with that by `trullekrul`, for the solution on the first nontrivial branch at $\lambda = 4$ (see (a)). Clearly, `e2rs` in (b) correctly identifies the triangles that are reasonable to refine, but refining the longest edges in Euclidean metric gives poor approximations at the 'boundary layers'. (c) shows a 'standard' refinement of (a) by `adapt_mesh` from `trullekrul` (with the given parameters from `troptions2D`), and (d) shows a refinement of (a) by `tradapt` with `sw=3`, i.e., without coarsening (and without swapping). This gives significantly more mesh points than (c), and our main purpose here is to illustrate the a-posteriori coarsening of (d) in (e,f), which gives a rather similar mesh as in (c).

Coarsening steps as in Fig. 1 from (d) to (e) are in particular useful for mesh adaptation during continuation. In Fig. 2 we continue the solution from Fig. 1(a) (at $\lambda = 4$ and $d = 0$) in d with mesh adaption each 5th continuation step, again comparing mesh adaptation by `trullekrul` with the old ad hoc adaptation by coarsening to the background mesh and then refining. The black branch in (a) belongs to the old option. The meshes and solutions (see (b) for two example plots) generally appear reasonable, but for larger d the interpolation down to the coarse background mesh and subsequent

Table 2: Control of `tradapt` via `sw` based binary coding with the 1st/2nd/3rd/4th bit switching on moving/refinement/coarsening/swapping, abbreviated as m/r/c/s, respectively. `sw=15` thus corresponds to the original `trullekrul adapt_mesh` behaviour.

<code>sw</code>	0	1	2	3	4	5	6	7	8	9	10	11	12	13	14	15
<code>action</code>	none	m	r	m,r	c	c,m	c,r	c,r,m	s	m,s	r,s	m,r,s	c,s	c,m,s	c,r,s	c,r,m,s

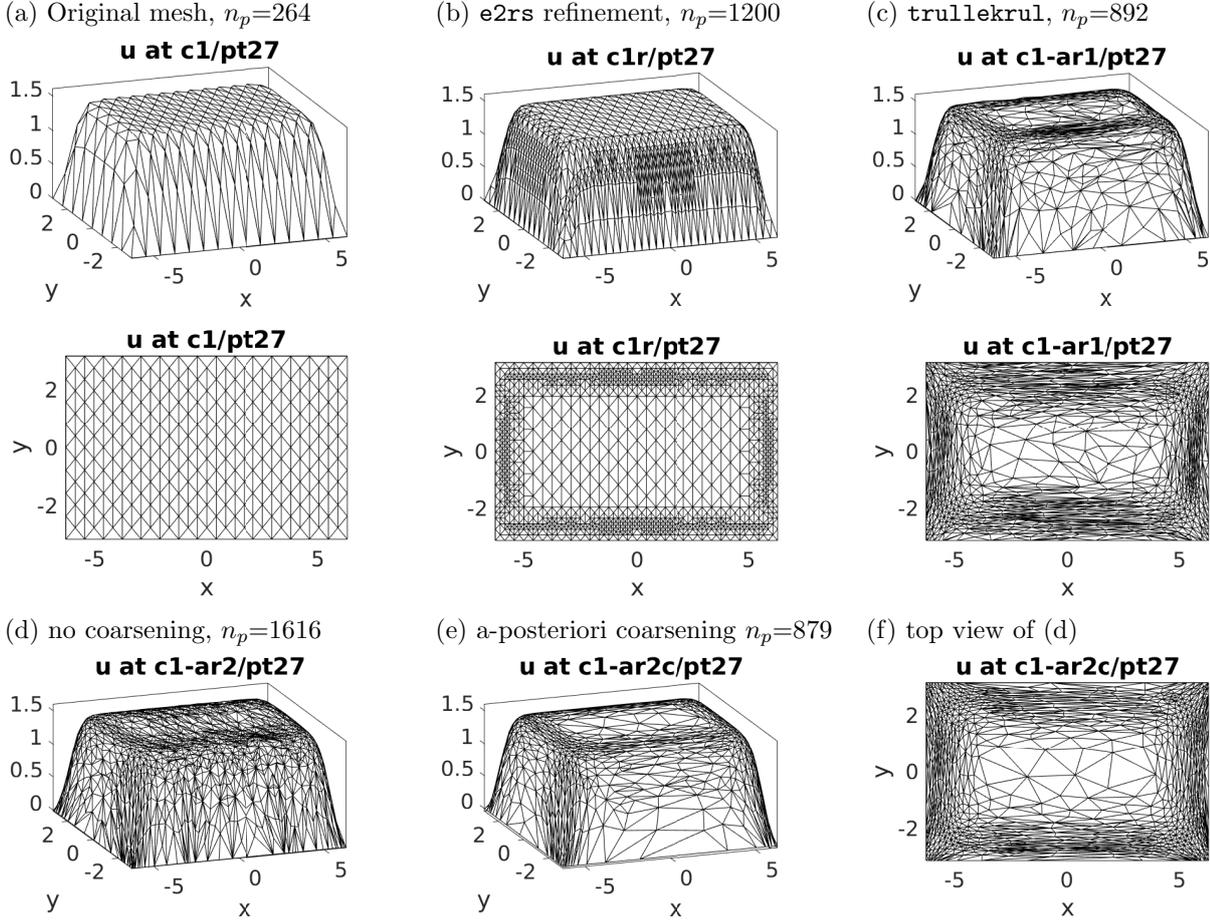


Figure 1: Selection of plots generated in `ac2D/cmds2.m`, illustrating different options for mesh-adaptation, see text for comments.

refinement become problematic, i.e., both appear somewhat underresolved at the $x = 2\pi$ boundary. A typical sign for such problems due to an inadequate background mesh are jumps in the branch data at adaption, that in (a) start to appear on the black branch for $d > 3$. The controlled coarsening–refine approach by `trullekrul` (red branch in (a)) is more robust in this respect. Of course, this is just one example, but it indicates a general result: if solutions during continuation develop boundary layers, or become strongly localized in some sense, use the `trullekrul` mesh adaptation. Listing 1 shows some pertinent commands for this for easy review.

```

45 % mesh-adaption during continuation, trullekrul
p=swiparf('c1','pt27','dia',4); p.sol.ds=0.1; p.nc.lammax=5; p.sw.trul=1;
op=troptions2D(); op.verbose=2; op.etafu=@etafua;
p.trop=op; % put trulle-options into p
p.nc.ngen=2; p.nc.amod=5; p=cont(p,30);
% mesh-adaption during continuation, trullekrul, with add. coarsening
p=swiparf('c1','pt27','dib',4); p.sol.ds=0.1; p.nc.lammax=5; p.sw.trul=1;
op=troptions2D(); op.verbose=2; op.etafu=@etafua;

```

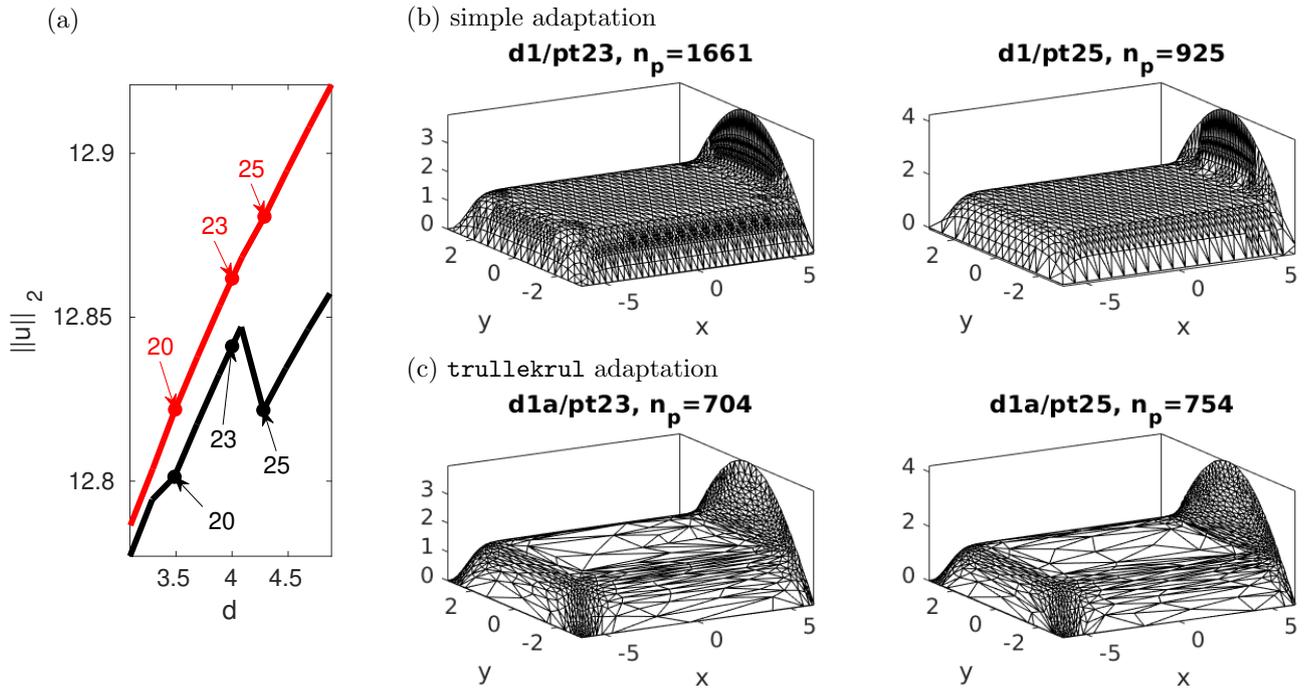


Figure 2: Further results from `cmds2`. Mesh adaptation every 5 steps of continuation in d , using `e2rs` and `trullekrul`, starting with the coarse mesh/solution from Fig. 1(a). See text for further comments.

```

50 p.trop=op; % put trulle options into p
   op.npb=500; % desired number of points after coarsening step
   op.sw=5; p.trcop=op; % put trulle coarsening options into p
   p.nc.ngen=2; p.nc.amod=5; p=cont(p,30);

```

Listing 1: Commands from `ac2D/cmds2.m` for continuation in d with mesh adaptation by `trullekrul`.

A wandering boundary spot. In a second example we consider (2) on the rectangle $\Omega = (-2\pi, 2\pi) \times (-\pi, \pi)$ with the BCs

$$\begin{cases} u = \exp(-(x - \xi)^2 - z^2) \text{ on } \Gamma_3 = \{y = \pi\}, \text{ parameter } \xi, \\ u = 0 \text{ on } \Gamma_1 = \{y = -\pi\}, \\ \partial_n u = 0 \text{ on } \partial\Omega \setminus (\Gamma_1 \cup \Gamma_3). \end{cases} \quad (7)$$

The purpose is to illustrate the mesh adaptation during continuation by considering a 'wandering spot' (upon continuation in ξ) on the top boundary. Additionally, this gives the opportunity to show how to put a parameter into the assembling of boundary values. The demo directory is `ac2Dwspot`, and Listing 2 shows the implementation of the rhs G , where we need to assemble the BCs in each step.

```

function r=sGws(p,u) % wandering boundary spot, parameter dependent BCs
2 par=u(p.nu+1:end); u=u(1:p.nu); xi=par(4); gr=p.pdeo.grid;
  bc1=gr.robinBC(0,0); bc3=gr.robinBC(1,0); % Neumann and DBC
  bcs=['exp(-(x-' mat2str(xi) ')'.^2)']; % parameter dependent BCs
  bc2=gr.robinBC(1,bcs); gr.makeBoundaryMatrix(bc3,bc1,bc2,bc1);
  [Q,Gbc,~,~]=p.pdeo.fem.assemb(gr); % the BC matrices
7 f=par(2)*u+u.^3-par(3)*u.^5; % the 'nonlinearity'
  r=par(1)*p.mat.K*u-p.mat.M*f+p.nc.sf*(Q*u-Gbc); % the residual

```

Listing 2: `ac2Dwspot/sGws.m`, G for the wandering spot example. Due to the ξ dependence, here we need to assemble the BCs G_{BC} in every call.

Fig. 3 (see also Listing 4) shows a continuation in ξ in the subcritical case ($\lambda = -0.25$), where the solutions are essentially characterized by the position of the boundary spot. For illustration we aim at rather coarse meshes, set `amod=5` (mesh-adaptation every 5th step), and allow for extra coarsening steps in `trullekrul`. The BD in (a) shows that this yields a reasonably smooth branch, and the sample plots in (b) that (as expected) the finest mesh may slightly lag behind the spot position (see in particular `pt16`), but otherwise the coarsening/refinement setup works very well.

```

20 % continue with trulle-adaptation with coarsening; -experiment with the params!
p=loadp('wsrc','pt0','wsada'); p=resetc(p); stansavefu(p); % reload pt and reset
p.nc.amod=5; p.nc.ngen=2; % p2p pars: adapt each amod-th step, in 2 iterations
p.trop.innerit=2; % resetting some trullekrul options (for testing)
p.trcop.npb=800; p.trcop.innerit=3; p.trcop.cmax=5; % resetting coarsening pars
p=cont(p,40); % run the continuation

```

Listing 3: Selection from `ac2Dwspot/cmds1.m`. Continuation in ξ , $\lambda = -0.25$ (subcritical case).

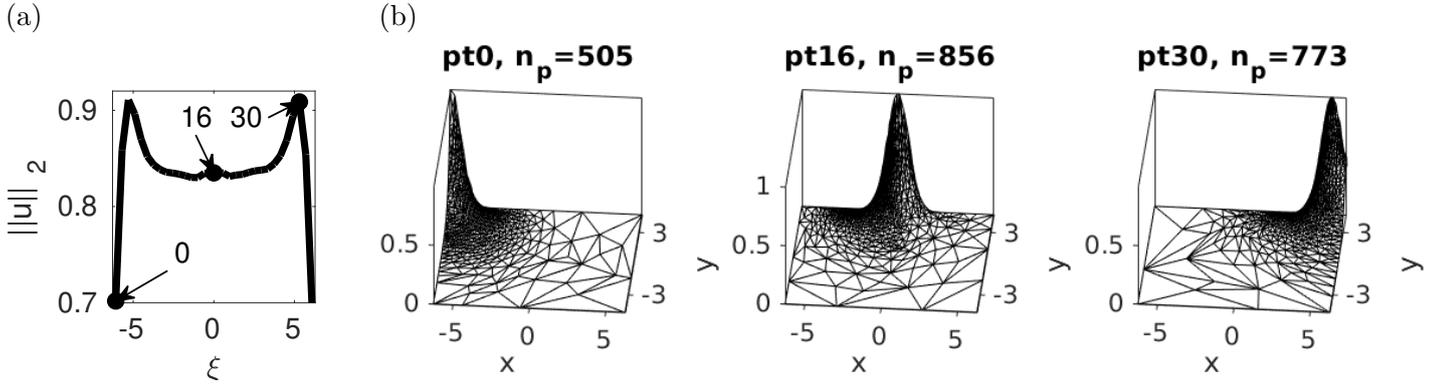


Figure 3: Example results from `cmds1.m` for (2) with the BCs 7, $(c, \lambda, \gamma) = (0.5, -0.25, 1)$, continuation in ξ , with mesh-adaptation every 5th step.

In `cmds2.m` we then continue in λ at fixed $\xi = 0$. The black branch in (a) initially corresponds to the 'trivial' branch on which solutions are as in Fig. 3 (specifically `pt16` at $\xi = 0$). Continuation in λ then yields an imperfect bifurcation to the primary unimodal branch in (a). Moreover, there are two BPs on the black branch, connected by the red branch. Altogether, the `trullekrul` mesh adaptation with `amod=10` yields robust results with still rather coarse meshes.

```

% continue soln at xi=0 in lambda, swiparf, then reset some pars
p=swiparf('wsada','pt16','lamc',2); p.nc.dsmax=0.02; p.sol.ds=0.01;
p.sw.foldcheck=0; p.nc.mu2=0.01; p.nc.foldtol=0.2; % reset foldtol (poor loc)
p.trcop.npb=1500; p.trop.etafu=@etafub; % allow finer meshes
5 p=cont(p,40); % go

```

Listing 4: Selection from `ac2Dwspot/cmds2.m`, continuation in λ , $\xi = 0$. We allow somewhat finer meshes, with `etafub = 10-6np`. The remainder of `cmds2` computes the bifurcating branch and then plots.

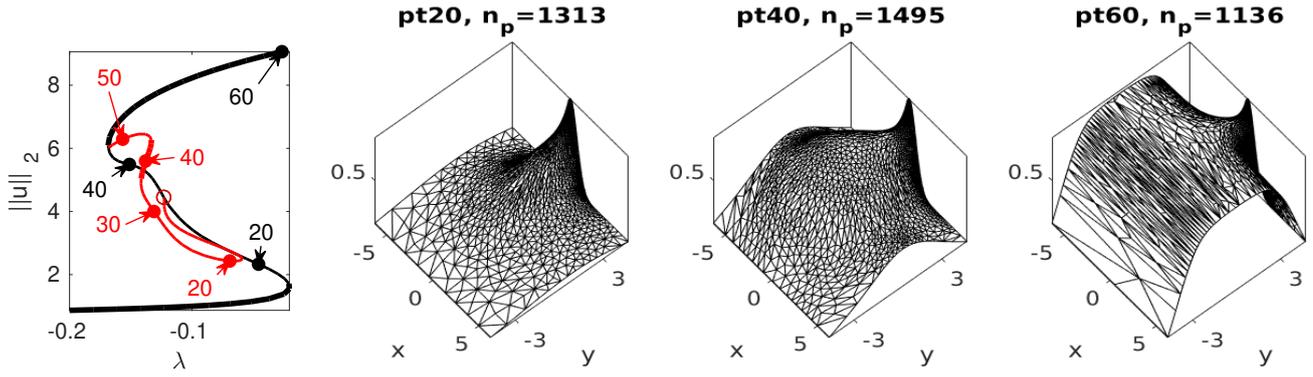
3.2 3D

In `ac3Dwspot` we consider analogous BCs as in (7), i.e.

$$G(u) := -c\Delta u - \lambda u - u^3 + \gamma u^5 = 0 \text{ in } \Omega, \quad (8a)$$

$$\begin{cases} u = \exp(-(x - \xi)^2 - z^2) \text{ on } \Gamma_3 = \{y = -3\pi/2\}, \text{ parameter } \xi, \\ u = 0 \text{ on } \Gamma_5 = \{y = 3\pi/2\}, \\ \partial_n u = 0 \text{ on } \partial\Omega \setminus (\Gamma_3 \cup \Gamma_5). \end{cases} \quad (8b)$$

(a) $\xi = 0$, continuation in λ and sample plots from black branch



(b) Sample plots from red branch

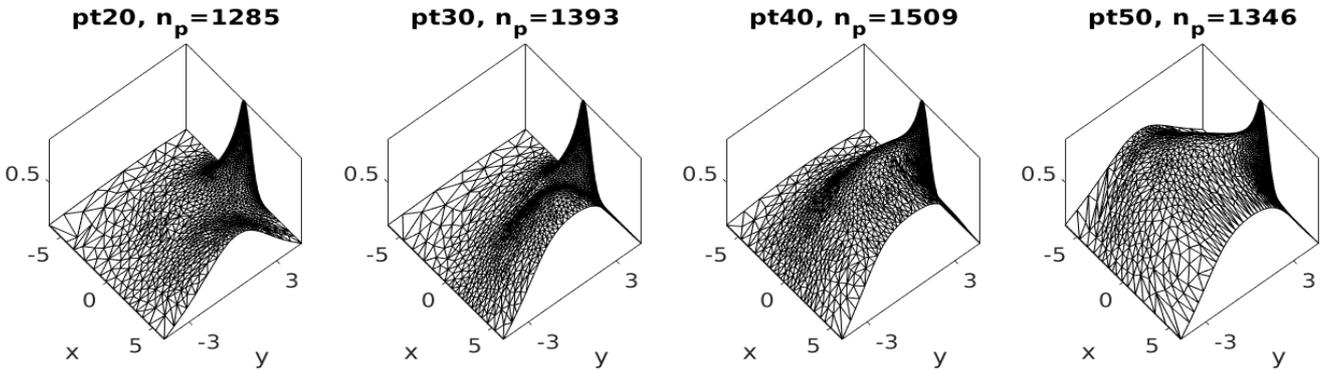


Figure 4: Example from `cmds2`. Continuing the primary solution branch in λ , fixed $\xi = 0$ yields an ‘imperfect bifurcation’ to the primary unimodal branch, and with an additional bifurcation to a branch with broken $x \mapsto -x$ symmetry, which, somewhat unexpectedly contains stable solutions, e.g., pt40.

For 3D cuboids as generated by `stanpdeo3D`, the order of the boundary faces is bottom, left, front, right, back, top, which leads to a straightforward modification of `sGws` from 2D to 3D. In `cmds1` we again we start with a continuation in ξ in the (here weakly) subcritical regime $\lambda = 0$, with the main results given in Fig. 5, using a very coarse initial mesh with $n_p = 3543$ to illustrate some effects and important settings for mesh-adaptation. Solutions on this coarse mesh look quite reasonable, see (a), and adaptive refinement (b) and subsequent coarsening (with `p.trcop.sw=5`, see Table 2) yield solutions which are visually the same, at least in the surface plots. However, for continuation on the fixed original coarse mesh from (a), for instance the L^2 -norm shows some unexpected fluctuations (red branch in (d)), e.g., for $\xi \in (0, 2)$. These are essentially due to the spot at a given ξ sitting on (close to) a mesh point, or in between two mesh points. The black branch in (d) is from continuation with mesh adaptation every 5th step, starting from the solution in (c), and setting `p.trcop.npb=3000` to coarsen a given mesh to at most 3000 mesh points before refinement. The two sample plots on the right of (d) are from this blue branch. Some main observations are:

1. The L^2 -norm on the adapted meshes is generally slightly smaller than on the (coarse) uniform mesh, and
2. The `trullekrul` coarsening (to less than 3000 mesh points) — refine approach yields nicely moving grids of small size with the finest meshing centered at the spots after each refinement.
3. Even though we only adapt every 5 steps, during which the spot moves to distance about 1 to 1.5 from the finest meshing, the black branch is reasonably smooth. However, if for instance we set `amod=10` (such that the spot moves further away before remeshing), then visible jumps occur in the L^2 -norm branch at each adaptation.

As in 2D, in `cmds2` we then switch to λ continuation, see Fig. 6. The basic behavior is as in 2D,

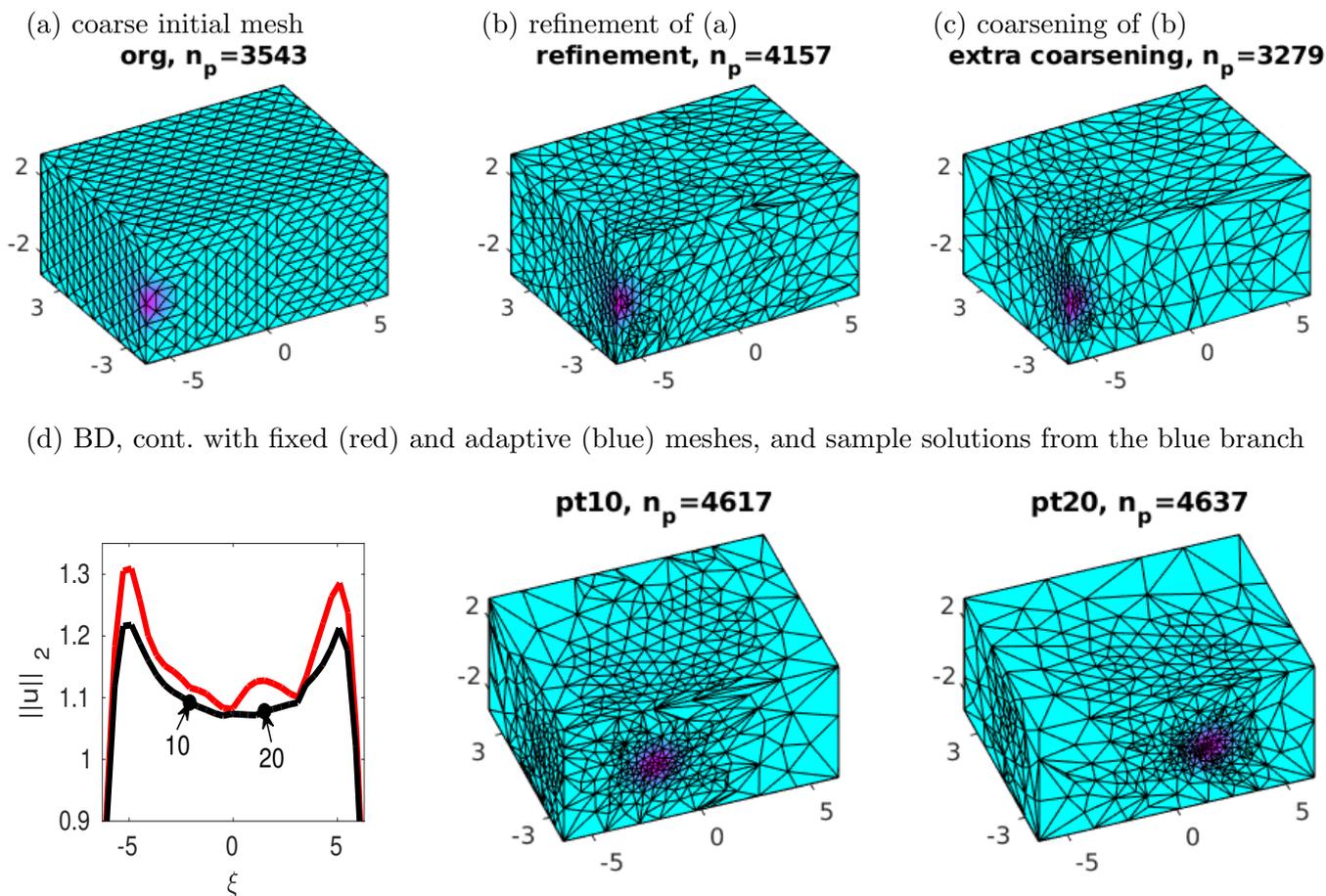


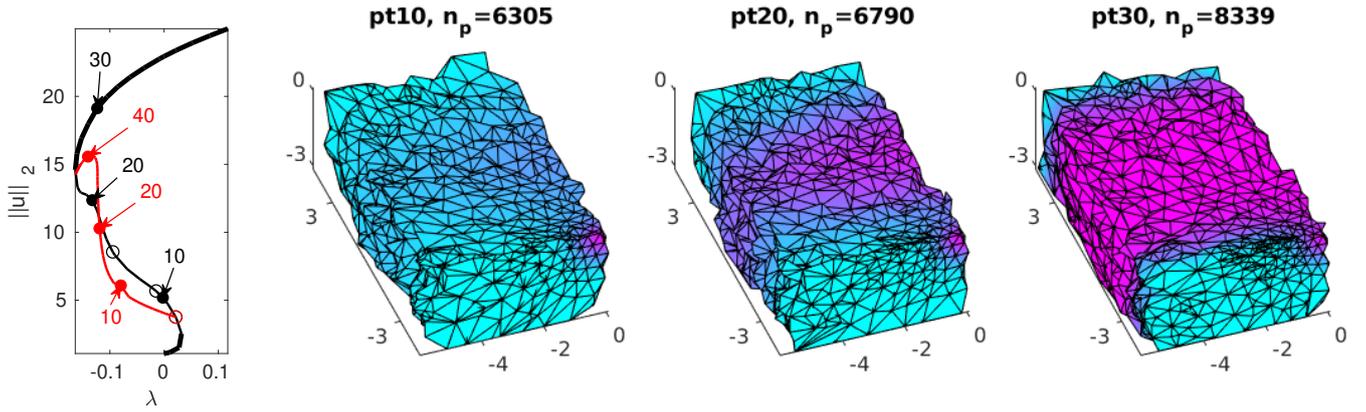
Figure 5: Selected plots from `ac3Dwspot/cmds1`. Colorscale in all plots from $u = 0$ to $u = 1$.

i.e., the black branch turns into the primary unimodal (+spot) branch by an imperfect bifurcation, and there are now four bifurcation points connected by branches with broken $x \rightarrow -x$ symmetry.

References

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(a) BD and sample plots (lower left quarter domain) from the primary λ branch



(b) Sample plots (lower half of domain) from the first bifurcating branch (red)

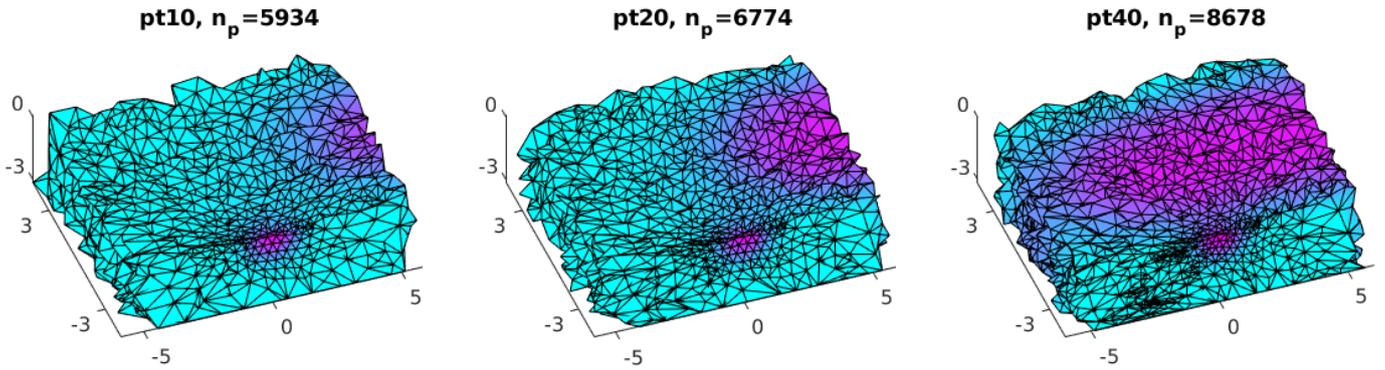


Figure 6: Selected plots from `ac3Dwspot/cmds2`. Colorscale in all solution plots from $u = 0$ to $u = 1$. Fixed $\xi = 0$, continuation in λ , yielding imperfect bifurcation to primary unimodal branch (black). At the first BP, a solution with broken $y \mapsto -y$ symmetry bifurcates, at the third BP the $z \mapsto -z$ symmetry is broken. The inner interfaces between $u = 0$ and $u = 1$ suggest a larger n_p . Some options used are: `amod=8`; `trop.qualP=2.25`; `trcop.npb=8000`.

[Uec19b] H. Uecker. Pattern formation with `pde2path` – a tutorial, 2019.

[Uec19c] H. Uecker. www.staff.uni-oldenburg.de/hannes.uecker/pde2path, 2019.

[UWR14] H. Uecker, D. Wetzel, and J.D.M. Rademacher. `pde2path` – a Matlab package for continuation and bifurcation in 2D elliptic systems. *NMTMA*, 7:58–106, 2014.