

# Rotated Slice Sampling for Efficient and Robust Estimation of DSGE Models

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### Introduction

- We investigate the performance of the rotated slice sampler first introduced by Planas Ratto Rossi (2015) (PRR2015) for the Bayesian estimation of medium/large scale DSGE models.
- We measure the relative performance of samplers based on inefficiency factors and we also test the rejection rates of the samplers against known distributions.
- We benchmark the rotated slice sampler together with the standard slice and the Metropolis-Hastings in the estimation of the Smets and Wouters (2003, 2007) model.
- the method is regularly used for the estimation of the European Commission's Global Multicountry model (Albonico et al., 2019).





### Main outcomes

- Unlike Metropolis, the Slice algorithm does not require any ad-hoc tuning nor any proposal distribution.
- The Rotated Slice extension boosts the efficiency of the sampler in case of highly correlated posterior distributions.
- A further extension of the rotated algorithm allows to dramatically increase the mixing properties of the slice sampler in successfully exploring the shape of complex multi-modal distributions.
- A parallel implementation of the algorithm provides accurate posterior draws with a computational cost comparable to the preliminary posterior maximization required by Metropolis ⇒ the slice algorithm strongly advisable in medium/large scale DSGE models.





# Bayesian inference of DSGE

Mostly using MCMC methods: obtaining draws from the the posterior distribution  $p(\theta, z|y)$  using the factorization:

$$p(\theta,z|y) = p(\theta|y)p(z|\theta,y)$$

 $\blacktriangleright \ \theta \sim p(\theta|y) \quad \Leftarrow \text{ our main concern}$ 

►  $z \sim p(z|\theta, y)$  off-line by a simulation smoother (e.g. Durbin and Koopman, 2002) using the Kalman filter.





# Sampling model parameters

Draws  $\theta \sim p(\theta|y)$  can be obtained in several ways:

- Random walk proposal Metropolis-Hastings (Dynare): easy to implement, faster than many other implementations, but sometimes inefficient.
- Slice sampler (Neal, 2003, also in Dynare): offers an avenue that we wish to explore carefully.

▶ ...





# Slice sampler's main features

Advantages:

- Only require  $f(\theta|y) \propto p(\theta|y)$  (like Metropolis);
- No proposal: does not require mode-Hessian or other info for defining proposal;
- Tuning parameters less important than Metropolis;
- Provides a framework for
  - adaptation
  - suppressing random walk behaviour (over-relaxation).

Disadvantages:

Not easy to generalize in the multivariate framework;

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Requires a large number of likelihood evaluations.





### In this talk

- The univariate slice sampler
- Some univariate test cases to mimic  $\theta_i \sim f(\theta_i | \theta_1, \cdots, \theta_{i-1}, \theta_{i+1}, \cdots, \theta_d, y)$ ,  $i = 1, 2, \cdots, d$
- Rotated univariate sampler correlated and multimodal distributions
- Test cases and DSGE applications
- Conclusions





# The idea of slice sampling

Introduce an *auxiliary variable*  $\gamma$  and construct  $p(\theta, \gamma)$  taking the marginal  $p(\theta)$  unchanged. Sampling from the joint  $p(\theta, \gamma)$  is not possible but ... draws from  $p(\theta)$  can be obtained iterating Gibbs updates on  $\gamma|\theta$  and  $\theta|\gamma$ :

sample  $\gamma$  given  $\theta$  from a uniform pdf over the set  $(0, f(\theta))$ 

Sample  $\theta$  given  $\gamma$  from a uniform pdf over  $S = \{\theta : \gamma < f(\theta)\}$ 





# Slice sampling in practice

Sampling  $\theta$  from a uniform over  $S = \{\theta : \gamma < f(\theta)\}$  is difficult to achieve exactly (perfect slice sampling). In practice:

▶ Position I = (L, R) around  $\theta^0$  at random that contains S as much as possible;

• Draw  $\theta$  from the set  $A = \{\theta : \theta \in S \cap I \text{ and } \Pr(I|\theta) = \Pr(I|\theta^0)\}$ 

Neal (2003) proposes some strategies:

(i) stepping out;

(ii) doubling;

(iii) random positioning, etc.





# Stepping out





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# The performance of the slice sampler

We start using a battery of univariate pdfs. By 1000 replications of sample size  $G=5000,\,\mathrm{we}$  measure

► NSE: 
$$Var(\frac{1}{G}\sum_{i=1}^{G}\theta^{i})^{1/2}$$
,  $i = 1, 2, \cdots, G$  (small)

▶  $\rho_1$ : the 1st order autocorrelation of the chain  $\theta^1, \cdots, \theta^G$  (close to 0)

► 
$$IF = 1 + 2\sum_{j=1}^{p} \omega_j \rho_j$$
,  $\omega_j$  the Parzen-weights (close to 1)

- the (average) number of calls to  $f(\theta)$  (small)
- ▶ the number of rejections of the Cramer-Von Mises (CVM) test (5%)





### Univariate test cases - Marron and Wand (1992)





### Univariate test cases - summary results

	RW-MH	Slice									
			Stepp	oing out			Doubling				
w		$\frac{1}{2}\sigma$ $3\sigma$ $10\sigma$ $100\sigma$				$\frac{1}{2}\sigma$	$3\sigma$	$10\sigma$	$100\sigma$		
NSE	3.77	2.75	1.92	1.83	1.79	2.30	2.02	1.85	1.80		
$ ho_1$	0.71	0.23	0.15	0.12	0.11	0.24	0.18	0.14	0.12		
IF	6.38	3.97	1.51	1.38	1.34	2.24	1.68	1.42	1.35		
N eval	1	9.42	6.09	6.61	9.72	23.31	14.65	9.47	10.00		
сум	0.59	0.22	0.13	0.11	0.11	0.20	0.15	0.12	0.12		
RE	1	4.1	1.42	1.42	2.04	7.44	3.72	2.12	2.12		





# Slice sampling for multivariate distributions

• Generalizing the stepping out procedure for a *d*-dimensional  $\theta$  (PRR2015):

- $\blacktriangleright$  need to evaluate  $f(\theta|y)$   $2^d$  -times to compute the vertices of the hypercube that approximates the Slice
  - $\Rightarrow$  parallelization as in Tibbits et al. (2011)?
- not easy to approximate the Slice in many dimensions by axis-aligned hypercubes  $\Rightarrow$  gradient of  $f(\theta)$ 
  - $\Rightarrow$  directional hypercubes for high-correlated parameters
- still difficult to solve the curse of dimensionality issue
- Here: Rotated univariate slice for high-correlated parameters as first proposed by PRR2015 and further developed here.





### Rotated univariate sampler

We use a rotated orthonormnal basis:

Let  $\Sigma \equiv Var(\theta|y)$ . Any rough estimate (e.g. burn-in period) will suffice.

By spectral decomposition  $\Sigma = A\Lambda A'$ .

- A, the eigenvectors of  $\Sigma$ , suggests in which direction to rotate the axes;
- the eigenvalues in  $\Lambda$  suggest the scale of the slices along each direction. We use  $W_i = 3\Lambda_{ii}^{1/2}$ .

This simple algorithm allows to increase the efficiency of the univariate sample for correlated multivariate distributions [ same cost as plain OAT for orthogonal ones].





Illustrations of the classical stepping out method by Neal (2003) (left) and the rotated stepping out method (right).





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# **DYNARE Implementation**

The rotated slice sampling is based on procedure with at least 2 steps:

- 1. burnin OAT, calling estimation with options
  - posterior\_sampling\_method='slice'
  - the parameter mh\_replic=50, mh\_blocks= ...
- 2. Rotated
  - posterior\_sampling\_method='slice'
  - posterior\_sampler\_options= ('rotated',1,'use\_mh\_covariance\_matrix',1)
  - load\_mh\_file, which is loading the variance-covariance matrix obtained in the previous step.
  - the parameter mh\_replic=50-1000, mh\_blocks= ...





### Multivariate test cases

We test the accuracy of some multivariate slice algorithms in four different examples:

1.  $\theta \sim N_d(0, I_d), d = 2, 5, 10$ 

 $\Rightarrow$  highlights issues even for well behaving distributions

- 2. Bivariate Mixture of 2 Normals (Chib Ramamurthy, 2010)  $\Rightarrow$  slow mixing of RW-MH even for small dimension; effectiveness of rotated slice
- 3.  $\theta \sim N_d(0, \rho 11' + (1 \rho)I_d), d = 10$   $\Rightarrow$  shows the effectiveness of rotated slice
- 4. Mixture of 4 Normals (Chib Ramamurthy, 2010) d = 12

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 $\Rightarrow$  shows the effectiveness of rotated slice





# Case 1. Gaussian - uncorrelated variables

 $\theta \sim N_d(0, I_d), d = 2, 5, 10$ 

	F	RW-MI	н		Slice					
				On	One-at-a-time					
d	2	5	10	2	5	10				
NSE	5.0	5.8	8.2	1.4	1.4	1.4				
$\rho_1$	.78	.90	0.95	.00	.00	.00				
IF	7.84	17.7	40.7	.95	.96	.95				
N eval	1	1	1	12	30	60				
сум	.83	.95	.99	.06	.07	.07				
RE	1	1	1	1.46	1.62	1.40				



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# Case 2. Chib Ramamurthy (2001)

$$\begin{split} \theta &\sim .99 \times N_2(\mu_1, \Sigma_1) + .01 \times N_2(\mu_2, \Sigma_2) \\ \text{where } \mu_1 &= (1, -1), \ \mu_2 = 6\mu_1, \ \Sigma_1 = 1.3I_2, \text{ and } \Sigma_2 = 0.05I_2. \end{split}$$

	RW-MH	Slice
		One-at-a-time
Max NSE	5.28	2.87
${\sf Max}\;\rho_1$	0.81	0.04
Max IF	12.94	11.9
N eval	1	11.99
сум	0.82	0.2
RE	1	11.02





# Case 2. Chib Ramamurthy (2010)





### Case 3. Gaussian - high correlated variables

	RW-MH	Slice				
		OAT	Rotated OAT			
Max NSE	10.4	11.3	1.8			
Max $ ho_1$	.98	.94	0.2			
Max IF	91	122	3.3			
N eval	1	71.2	61			
CVM	1	.96	0.1			
RE	1	95.6	2.2			





# Slice sampler

- Rule of thumb for the tuning parameter  $W \simeq 3\sigma$ ;
- Stepping out procedure generally outperforms the doubling one (and other not reported here);
- Univariate slice sampling more appealing than multivariate one when variables are loosely correlated;
- Rotated univariate sampler works well with highly-correlated variables and multi-modal problems.
- How bout medium-large scale problems?





# DSGE examples

- Smets Wouters (2003, 2007)
- EC Global Multicountry model (Albonico et al., 2019);

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An Schorfheide (2007);





# Smets Wouters (2003)





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# Smets Wouters (2003, 2007)

#### SW2003 32 parameters

SW2007 36 parameters

	RW-MH	Slice		RW-MH	Slice	
		Rotated OAT			Rotated OAT	
IF	110	3.2	IF	110	3.3	
N eval	1	183	N eval	1	210	
RE	1	5.37	RE	1	6.3	





# Modified Implementation: go parallel

Once chains properly start, not clear trade-off between slice and MH. BUT: can slice be run at the same cost of optimizer??

- 1. burnin OAT, calling estimation with options
  - > posterior\_sampling\_method='slice'
  - ▶ the parameter mh\_replic=50  $\rightarrow$  30-35, mh\_blocks=250
- 2. Rotated
  - posterior\_sampling\_method='slice'
  - posterior\_sampler\_options= ('rotated',1,'use\_mh\_covariance\_matrix',1)
  - load\_mh\_file, which is loading the variance-covariance matrix obtained in the previous step.
  - ▶ the parameter mh\_replic=100-1000  $\rightarrow$  15-20, mh\_blocks=250





# Q: How short can be slice/MH chains?

- We test the performance of the slice sampler and MH for estimating the parameters of a SW2003-like posterior (approximated with a truncated normal).
- We apply the multivariate version of the Cramer test proposed by Baringhaus and Franz (2004)
- We explore the posterior of the model through samples of 250 points, obtained as the last element of 250 parallel chains (each point has 32 parameters/dimension)
- We repeat the experiment 30 times to get the distribution of the p-value of the test (each point has 32 parameters/dimension)





### Slice sampler performance after rotation - non truncated normal

We give the p-values of the Cramer test at different poinst of the chain, during burnin and after rotation. Null of the test: the sample comes from the known distribution of the posterior.

SLICE SAM	PLER CRAME	R TEST - NON	TRUNCATE	NORMAL												
	BURNIN						SAMPLER AFTER ROTATION									
	1	10	20	30	40	50	150	250	350	450	550	650	750	850	950	1050
μ	0.00	0.00	0.00	0.02	0.34	0.45	0.57	0.48	0.51	0.49	0.56	0.54	0.43	0.56	0.53	0.56
σ	0.00	0.00	0.00	0.02	0.21	0.29	0.26	0.34	0.27	0.27	0.28	0.29	0.24	0.28	0.30	0.27
#<0.05	30	30	30	27	1	3	1	3	0	0	0	1	1	0	1	1
%<0.05	100.0%	100.0%	100.0%	90.0%	3.3%	10.0%	3.3%	10.0%	0.0%	0.0%	0.0%	3.3%	3.3%	0.0%	3.3%	3.3%
SLICE SAM	PLER CRAME	R TEST - TRU	NCATED NO	RMAL												
			BUR	NIN			SAMPLER AFTER ROTATION									
	1	10	20	30	40	50	150	250	350	450	550	650	750	850	950	1050
μ	0.00	0.00	0.00	0.01	0.46	0.45	0.50	0.46	0.53	0.48	0.46	0.56	0.47	0.49	0.39	0.51
σ	0.00	0.00	0.00	0.01	0.23	0.29	0.26	0.30	0.27	0.28	0.29	0.22	0.28	0.34	0.30	0.30
#<0.05	30	30	30	30	2	2	0	3	0	2	3	0	2	5	2	0
% < 0.05	100.0%	100.0%	100.0%	100.0%	6.7%	6.7%	0.0%	10.0%	0.0%	6.7%	10.0%	0.0%	6.7%	16.7%	6.7%	0.0%





Adaptive Slice sampler performance before/after rotation - truncated normal

OAT length: 30-35 iterations; rotated length: 15-20 iterations.







### Metropolis Hastings - truncated normal

We randomize the p-values of the Cramer test at different point of the chain. Null of the test: the sample comes from the known distribution of the posterior.







# A: How short can be slice/MH chains?

- A few tens of slice iterations (without the fixed cost of optimizer)
- A few hundreds of MH iterations (*after* the optimization step)

are sufficient to get good iid sample from a lot of parallel chains

- Rotated slice iterations can start after about 20-30 iterations, reducing computational cost.
- How about overhead cost of optimization for MH?





### Computational cost of slice.

One slice iteration uses on average 6-8 function evaluation per parameter. So, for 50 iterations, the total number of function evaluations N for n parameters can estimated as:

 $N=50\times7\times n$ 

Note. Never set the 'usual' huge number of MCMC iterations applied for MH. Slice will last forever for such a big number!





### Computational cost: slice vs. optimizer

Number of function evaluations to run 50 slice iterations vs optimizer. Across different tests, optimizer may last more than the 50 slice iterations. Optimization becomes more difficult:

- when posterior is more concentrated w.r.t. prior
- presence of cliffs (e.g. no solution, B-K violations etc.)

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▶ large number of parameters *n*.







### Computational cost: slice vs. optimizer. n = 32 vs n = 192

Number of function evaluations to run 50 slice iterations vs optimizer. Increasing the size of the problem, cost of optimizer increases w.r.t. the 50 slice iterations.





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# Rotated univariate sampler: multimodal problems

The same idea can also be used to improve mixing and efficiency of the sampler in the case of multi-modal problems. As in Chib Ramamurthy (2010) rely on assumption that we know the location of the different local optima:

- 1. identify (at least some of) the multiple local optima characterizing the posterior distribution;
- 2. perform slice sampling along rotated axes, parallel to lines joining different local optima





# Case 2. Chib Ramamurthy (2010) REVISITED

$$\begin{split} \theta &\sim .99 \times N_2(\mu_1, \Sigma_1) + .01 \times N_2(\mu_2, \Sigma_2) \\ \text{where } \mu_1 &= (1, -1), \ \mu_2 = \frac{8\mu_1}{1}, \ \Sigma_1 = 1.3I_2 \text{, and } \Sigma_2 = 0.05I_2. \end{split}$$





# Case 4. Chib Ramamurthy (2010)

Mixture of 4 normals in 12 dimensions with probability  $p_j = [0.75, 0.05, 0.15, 0.05]$ 





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# Case 4. Chib Ramamurthy (2010)

Mixture of 4 normals in 12 dimensions with probability  $p_j = [0.75, 0.05, 0.15, 0.05]$ . Excerpt from Chib Ramamurthy:







# Case 4. Chib Ramamurthy (2010): go parallel

Parallel implementation of mixture of 4 normals in 12 dimensions with probability  $p_j = [0.75, 0.05, 0.15, 0.05]$ . It easily finds the four modes, but misses relative weight.





# Case 4. Chib Ramamurthy (2010): go parallel

Compute the marginal likelihood for the four optima:  $\hat{p}_j = [0.73, 0.05, 0.14, 0.05]$ . Re-weight estimated densities accordingly:





# An Schorfheide (2007)

- multimodal posterior distribution;
- although 'low' mode has very low probability (0.005%), MH or standard slice samplers incapable to jump to high density region
- rotated slice sampler can deal with this (likewise Chib Ramamurthy, 2010, with Tailored randomized block MCMC)





# An Schorfheide (2007)

Univariate slice sampler:



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# An Schorfheide (2007)

Rotated univariate slice sampler:





# An Schorfheide (2007): go parallel

- perform many short chains as usual: very easily get to the two modes.
- Compute the relative marginal likelihood for the two optima:  $\hat{p}_{2,1} = 0.03\%$ [relative likelihood at mode:  $\hat{L}_{2,1} = 0.005\%$ ]
- Re-weight estimated densities accordingly...





### Conclusions

Slice sampler has appealing properties for usage with DSGE models:

- does not require optimization to compute the mode and the Hessian;
- small IF and short chains: good to get a few 'good' points in the posterior space, with a limited initial budget of function evaluations;
- rotated slice makes the method much more efficient, even at short number of iterations;
- available in DYNARE.

[Note. Optimizer 5 of DYNARE is based on the same principle, with Gibbs steps.]





### Conclusions

Many short parallel chains are to be preferred to long chains:

- provides iid samples;
- applies to any kind of sampler MH/slice etc.
- applies to large problems;
- short slice chains (50 iterations) are often faster than the optimizer needed to start MH;
- effective also for multimodal problems.





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