



Transdimensional Markov Chain Monte Carlo Methods

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Motivation for Different Inversion Technique

- Inversion techniques typically provide a single best-fit model while the fit of other models may be only slightly worse.
- Inversion algorithms often get stuck in local minima.
- Non-uniqueness is often uncharacterized.
- Dimensionality (e.g. number of layers) may be unknown prior to inversion.
- Uncertainty analysis is desired on individual features within a result.

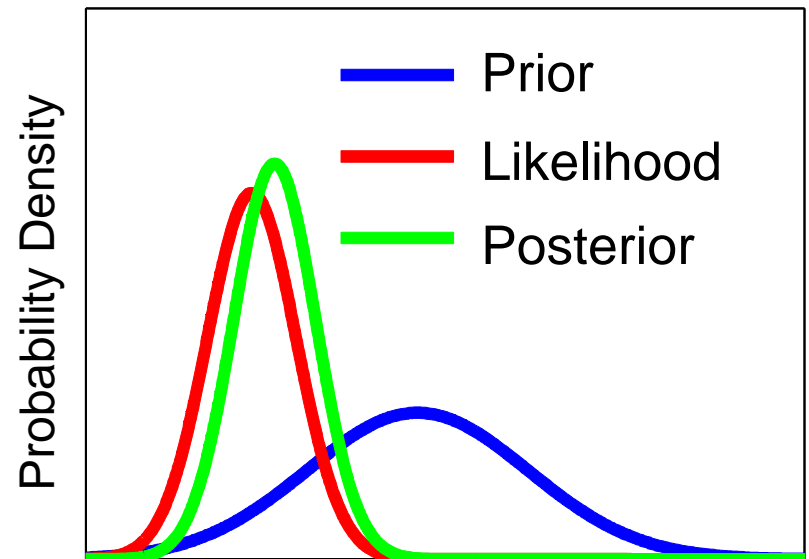
Bayes' Theorem

Incorporates prior knowledge and data to obtain posterior probability density function (PDF):

$$p(\mathbf{m}|\mathbf{d}) = \frac{p(\mathbf{d}|\mathbf{m})p(\mathbf{m})}{p(\mathbf{d})}$$

Labels for the equation:

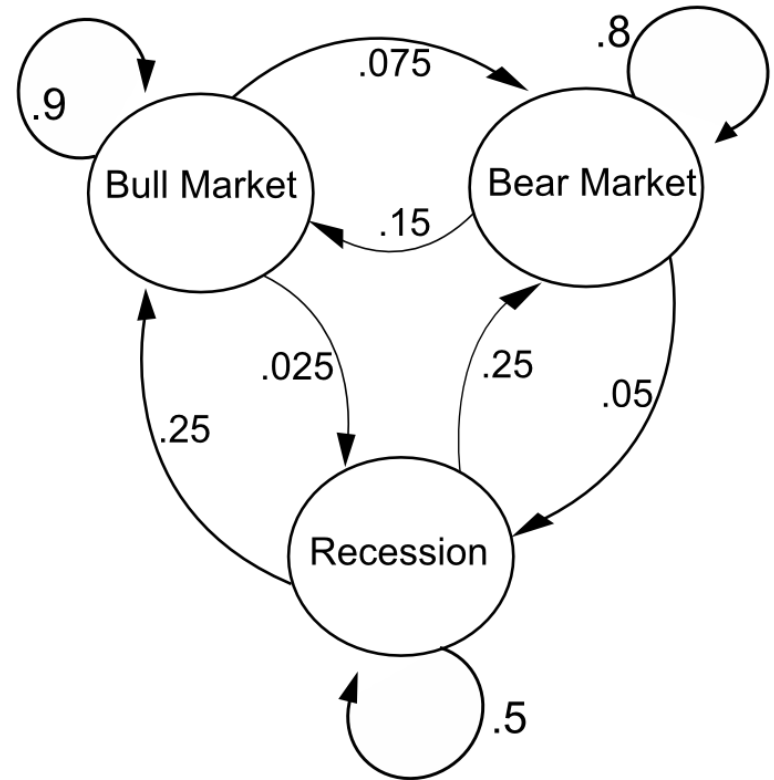
- Likelihood PDF (points to $p(\mathbf{d}|\mathbf{m})$)
- Prior PDF (points to $p(\mathbf{m})$)
- Posterior PDF (points to $p(\mathbf{m}|\mathbf{d})$)
- Evidence (points to $p(\mathbf{d})$)



What *is* Markov Chain Monte Carlo?

Markov chain: a process in which the next state only depends on the current state

Monte Carlo: Using random numbers to estimate properties of a solution



Example of 3-state Markov chain. Courtesy of Gareth Jones.

Metropolis-Hastings Algorithm

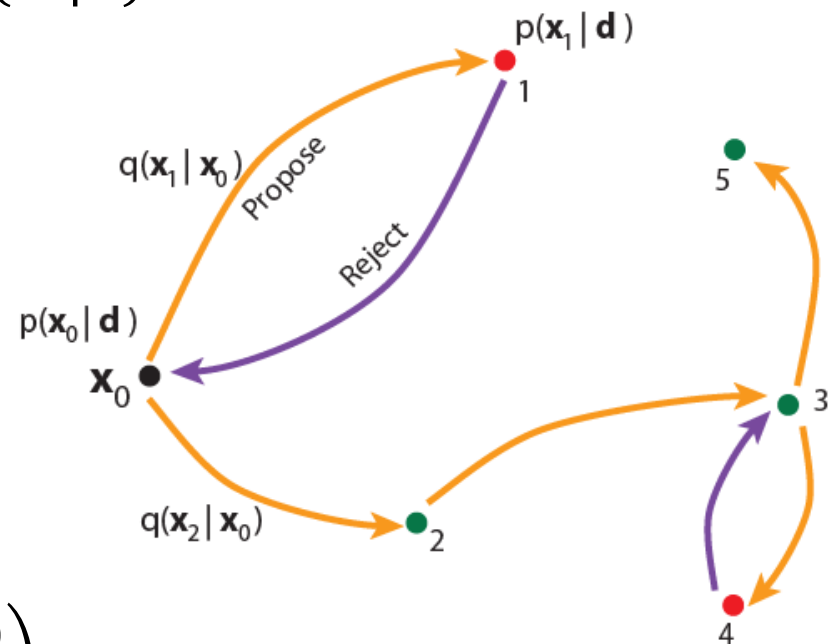
Tests new set of model parameters, m' , in order to sample the posterior probability distribution, $p(m|d)$.

Proposal distribution for m'
given current model, m :

$$q(m'|m)$$

Acceptance probability:

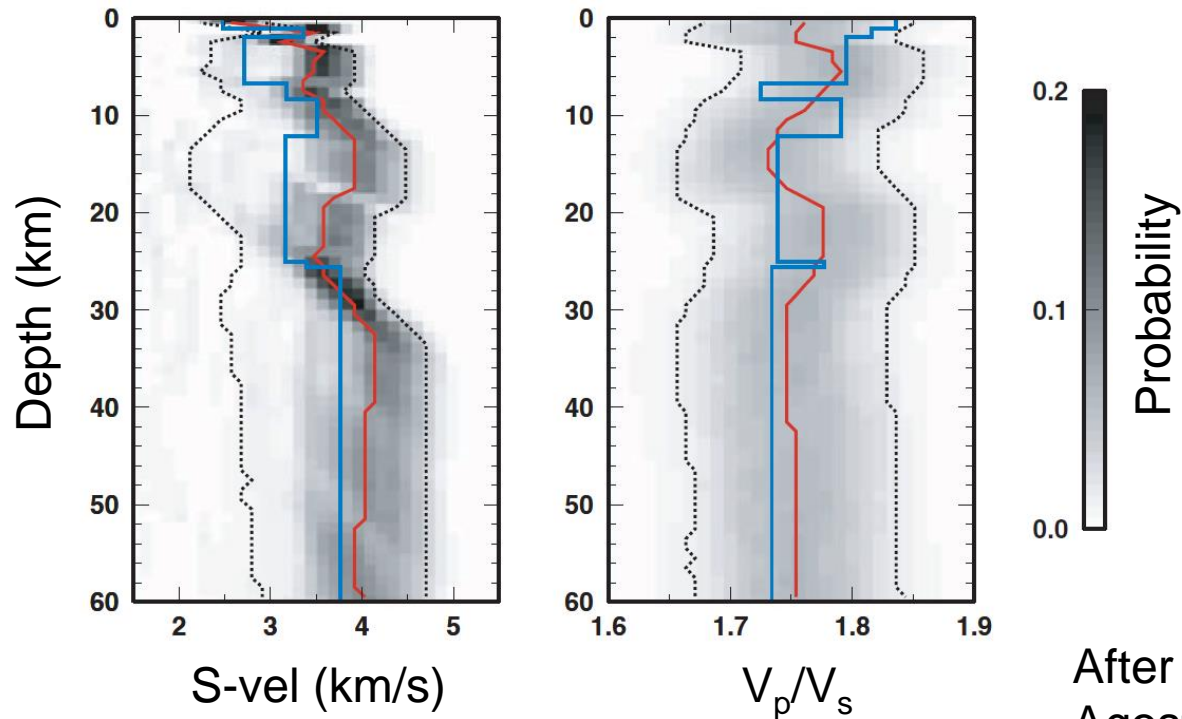
$$\alpha = \min \left(1, \frac{p(m')p(d|m')q(m|m')}{p(m)p(d|m)q(m'|m)} \right)$$



Example of progression of states. After Sambridge (2010).

MCMC Results

After Markov chain has converged, models are binned and the ratio of models in a given bin is proportional to posterior probability in that bin



After Piana
Agostinetti and
Malinverno (2010)

Reversible-Jump (Transdimensional) MCMC

Allows for jumps between parameter spaces of different dimensions

Acceptance probability changes from

$$\alpha = \min \left(1, \frac{p(\mathbf{m}')p(\mathbf{d}|\mathbf{m}')q(\mathbf{m}|\mathbf{m}')}{p(\mathbf{m})p(\mathbf{d}|\mathbf{m})q(\mathbf{m}'|\mathbf{m})} \right)$$

To

$$\alpha = \min \left(1, \frac{p(\mathbf{m}')p(\mathbf{d}|\mathbf{m}')q(\mathbf{m}|\mathbf{m}')}{p(\mathbf{m})p(\mathbf{d}|\mathbf{m})q(\mathbf{m}'|\mathbf{m})} |\mathbf{J}| \right)$$

Jumps between parameter spaces are commonly done using Birth-Death MCMC

RJMCMC for Receiver Function Deconvolution

- Receiver functions use P to S or S to P conversions from teleseismic waves to infer velocity structure
- Assumed relation between Parent (P) and Daughter (D) waveforms:

$$D = P * G + \varepsilon$$

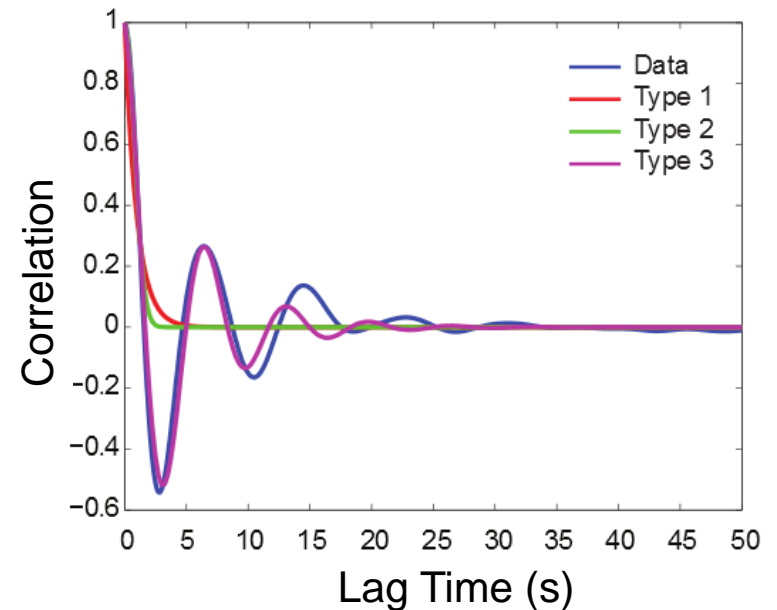
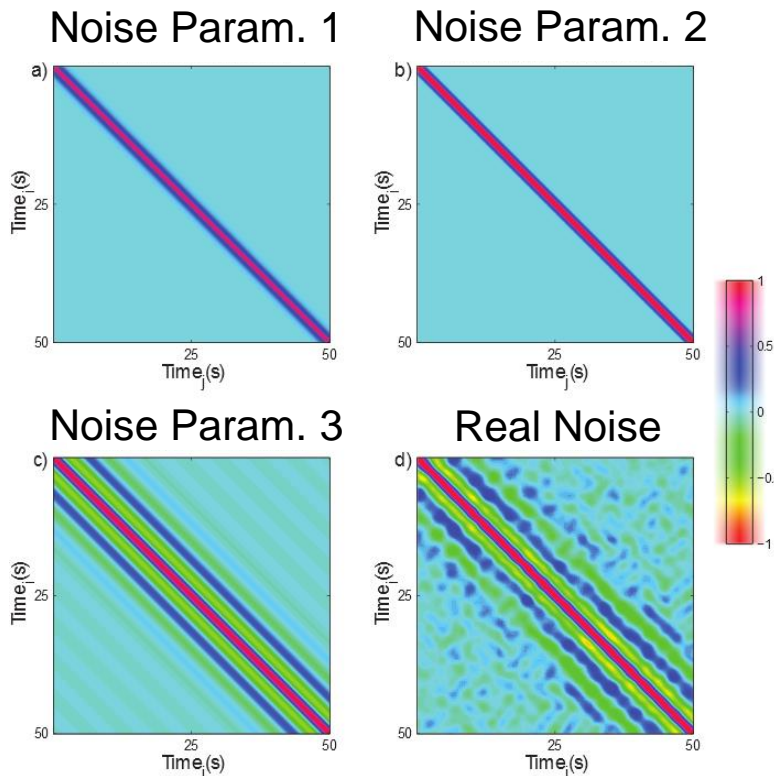
- Receiver function, G, is parameterized as an unknown number of Gaussians at unknown lag times with unknown widths and amplitudes.
- Likelihood of observed daughter waveform given G:

$$p(D|G) = \frac{1}{\sqrt{(2\pi)^n |C_D|}} e^{-(P*G-D)^T C_D^{-1} (P*G-D)}$$

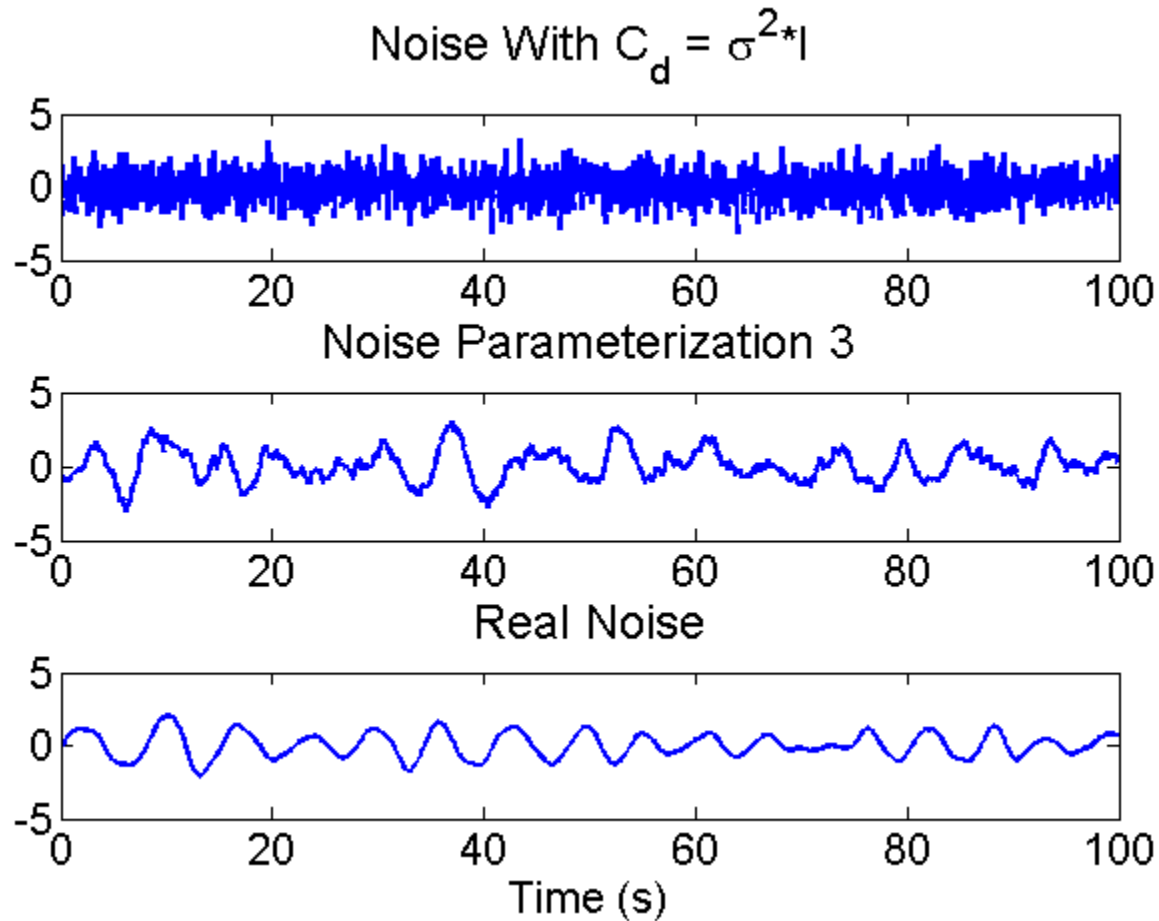
Noise Parameterization

We parameterize the noise covariance matrix with hyperparameters σ , λ , and ω_0 :

$$C_{D_{ij}} = \sigma^2 e^{-\lambda|t_j - t_i|} \cos(\omega_0 \lambda |t_j - t_i|)$$



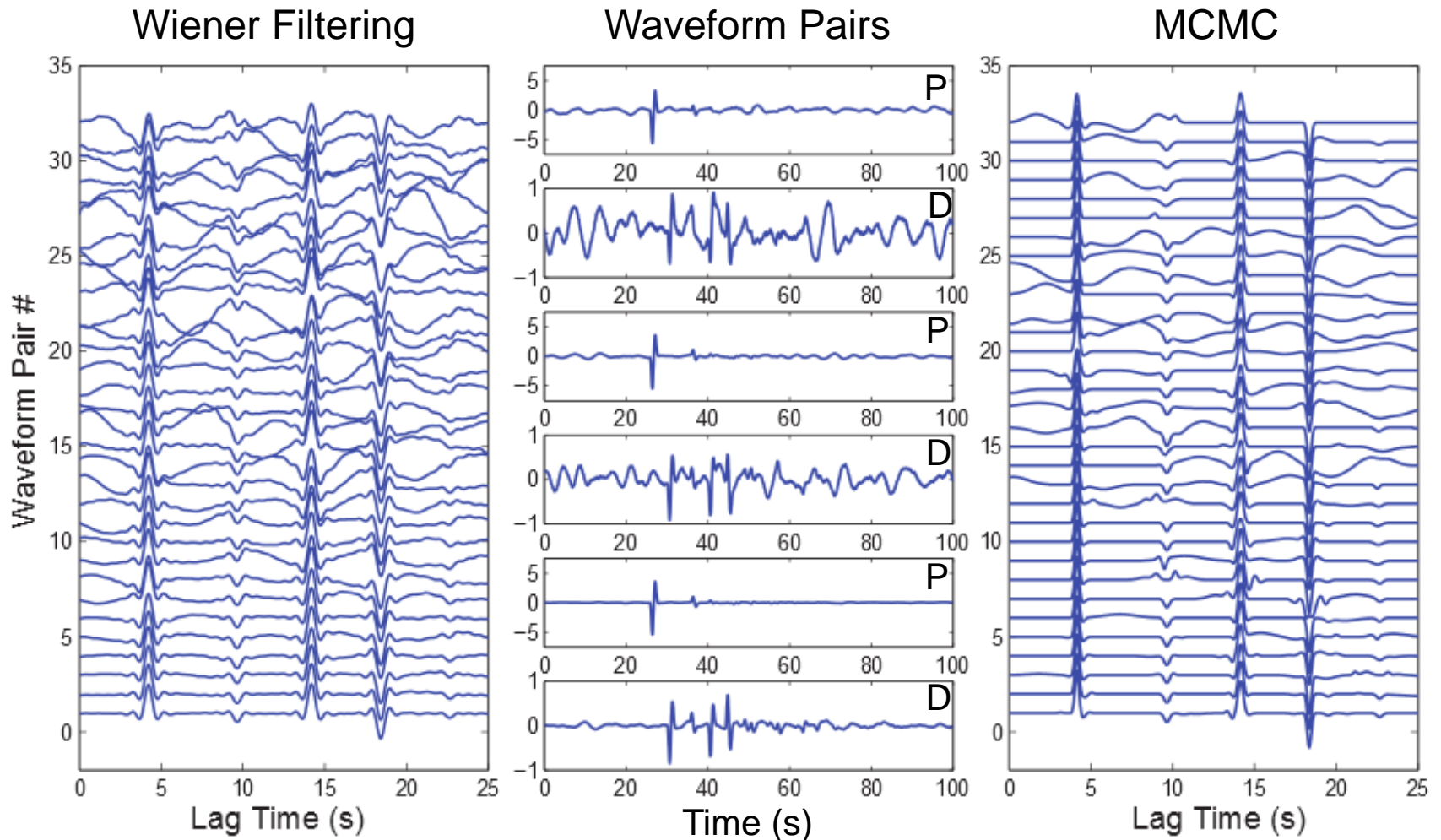
Why Correlation Matters



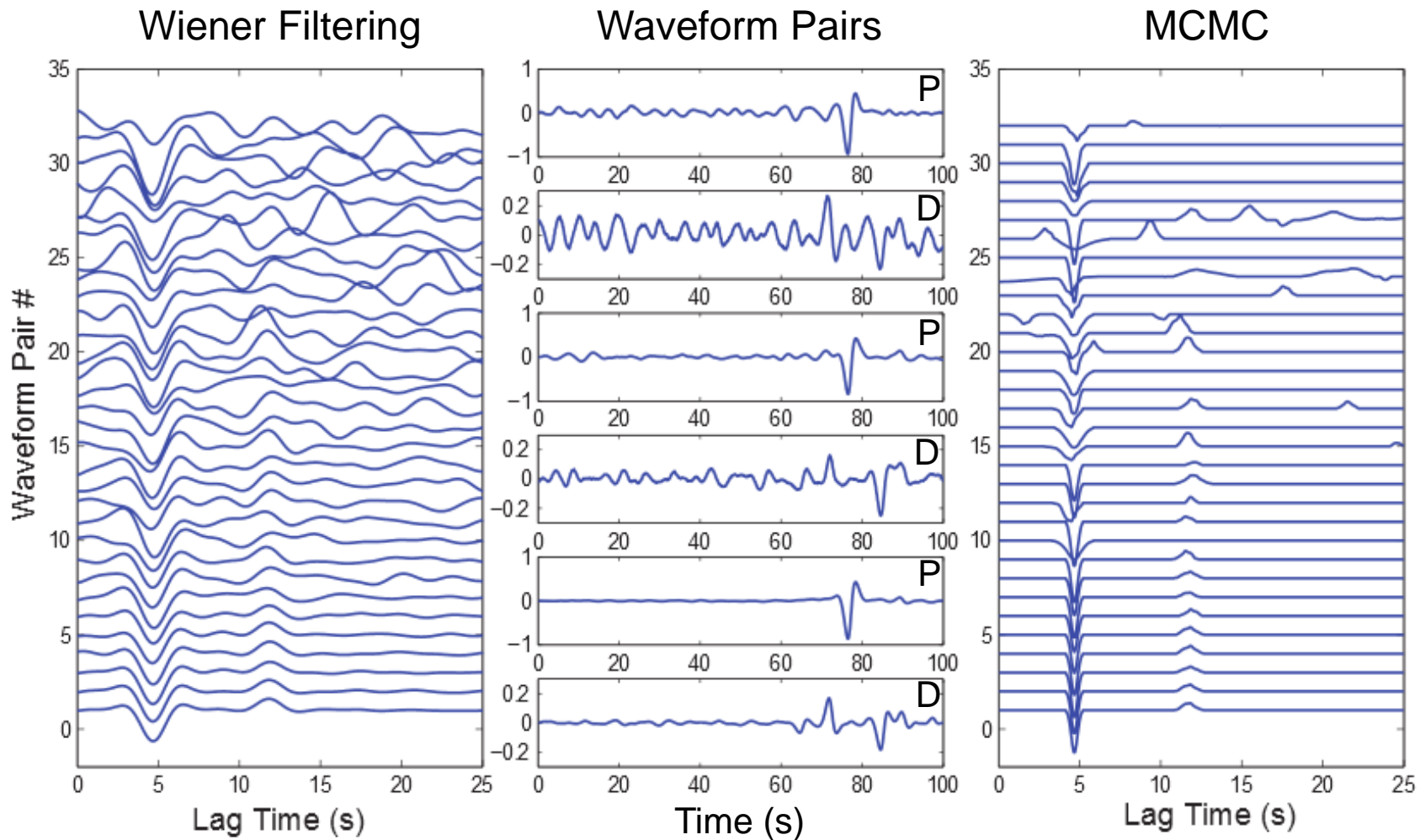
Process

- Start with initial model, G , of no Gaussians.
- Convolve P with G (forward model) and use to calculate likelihood.
- Propose new model by either “birthing” or “killing” a Gaussian, changing an existing Gaussian’s amplitude, width, or location, or changing a noise hyperparameter.
- Calculate forward model, likelihood, and acceptance probability of new model. If acceptance probability is greater than a random number from 0 to 1, accept the model. Otherwise, reject it.
- Repeat proposal and accept/reject process until convergence and then continue, saving models afterwards.

PS MCMC results



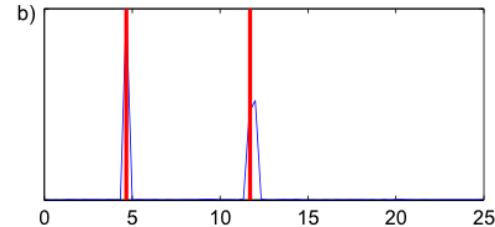
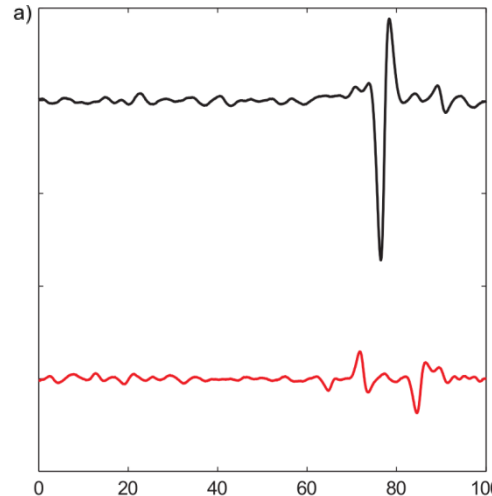
SP MCMC results



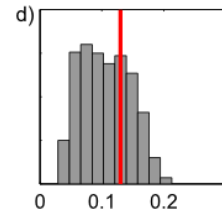
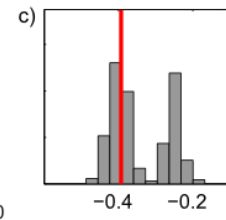
SP Result Analysis

Parent & Daughter Waveforms

Location

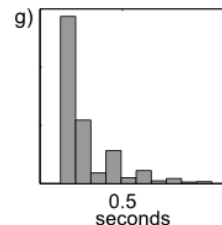
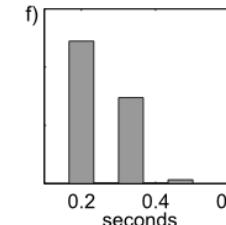
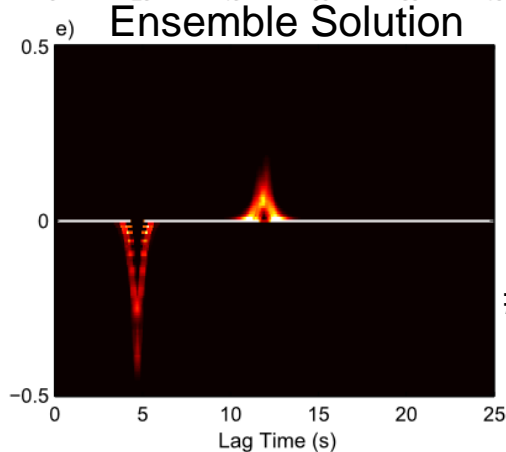


1: Amplitudes 2: Amplitudes



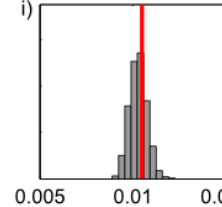
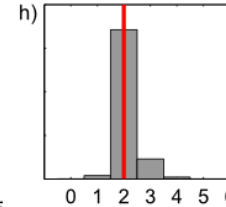
1: Widths

2: Widths



of Gaussians

Noise σ

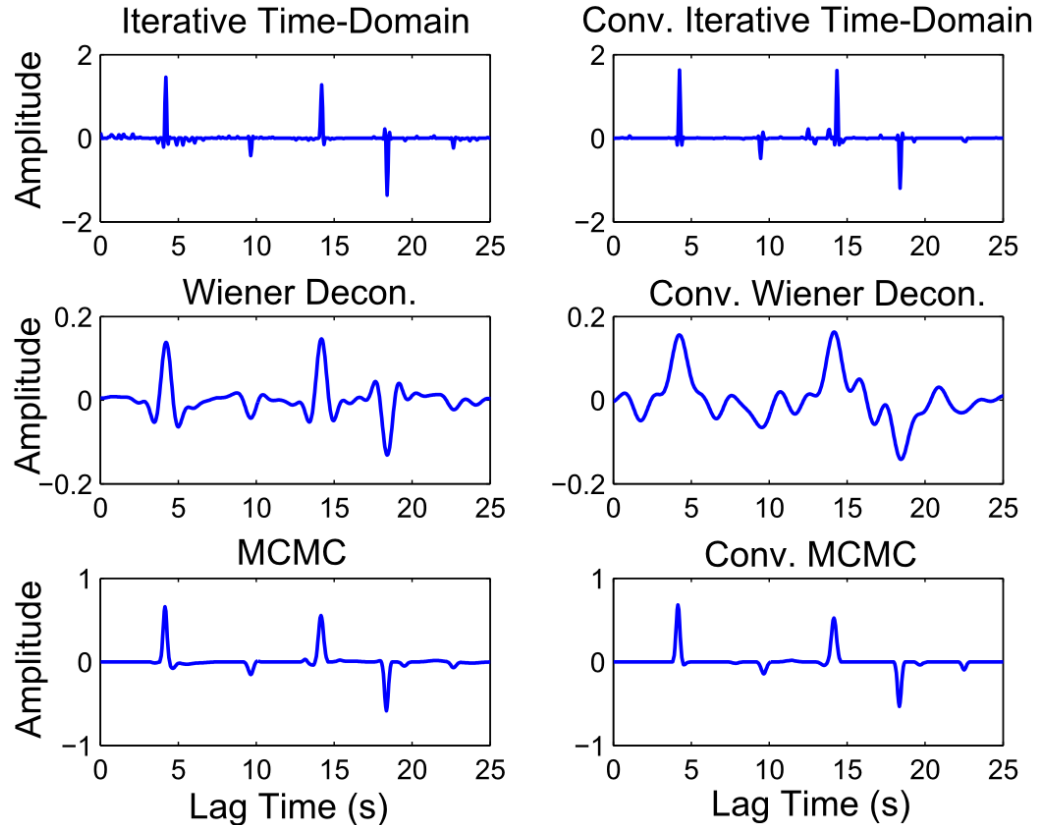


Low High
0 0.05 0.1 0.15
Posterior Probability

Spectral Hole Test

Solve for G when using STF with spectral holes:

$$(D * STF) = (P * STF) * G + \varepsilon$$



Model Comparisons

Can compare hypotheses H which may be parameterizations or forward models:

$$p(\mathbf{m}|\mathbf{d}, H) = \frac{p(\mathbf{d}|\mathbf{m}, H)p(\mathbf{m}|H)}{p(\mathbf{d}|H)}$$

With evidence, $p(\mathbf{d}|H)$:

$$p(\mathbf{d}|H) = \int p(\mathbf{d}|\mathbf{m}, H)p(\mathbf{m}|H)d\mathbf{m}$$

Point approximations using evidence:

$$AIC = -2\log[p(\mathbf{d}|\mathbf{m}_{mle})] + 2k$$

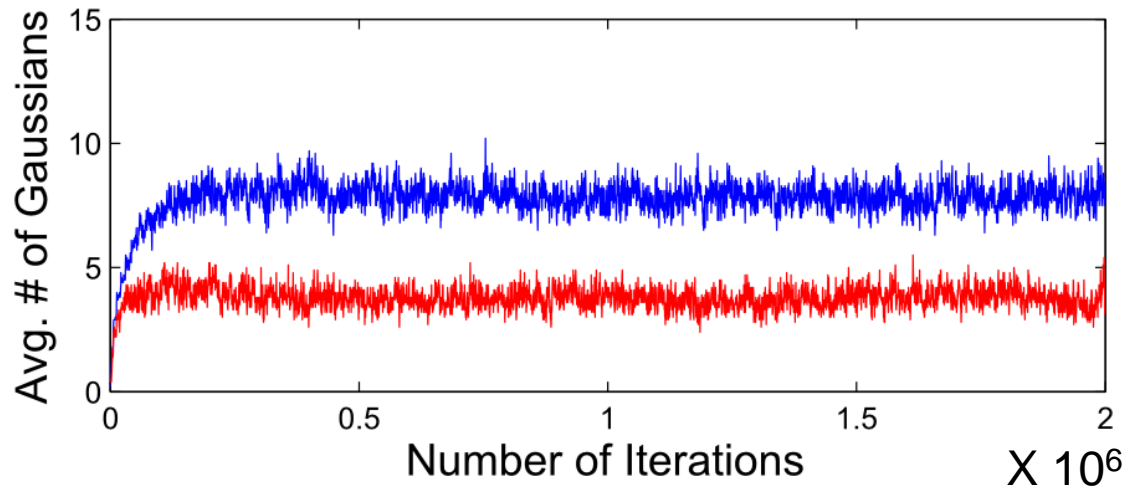
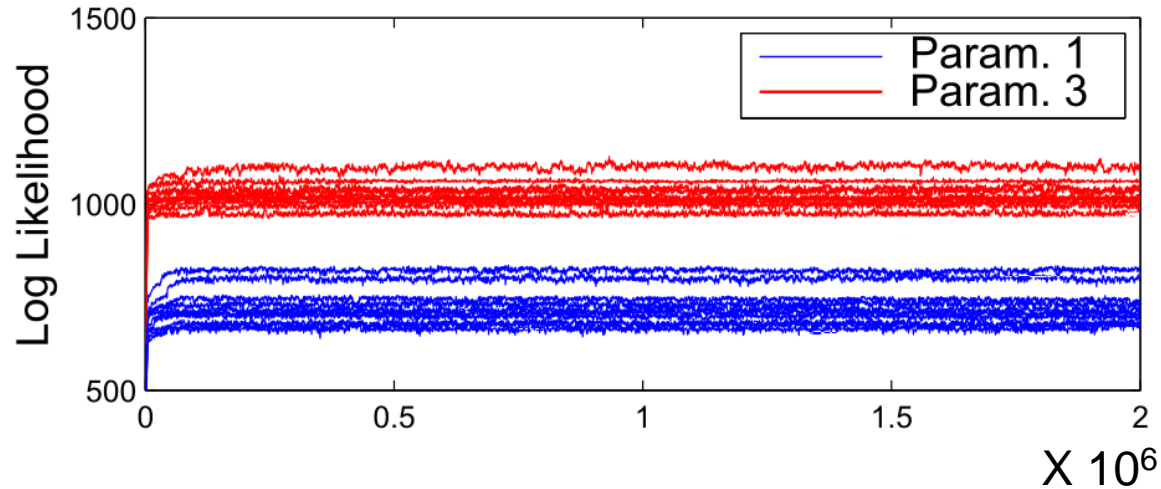
$$BIC = -2\log[p(\mathbf{d}|\mathbf{m}_{mle})] + k\log(n)$$

$$DIC = E^\theta[-2\log[p(\mathbf{d}|\mathbf{m})] + k_{eff}]$$

Where \mathbf{m}_{mle} is maximum likelihood estimate of the model, k is number of parameters, and n is sample size

Noise Parameterization 1 vs. 3

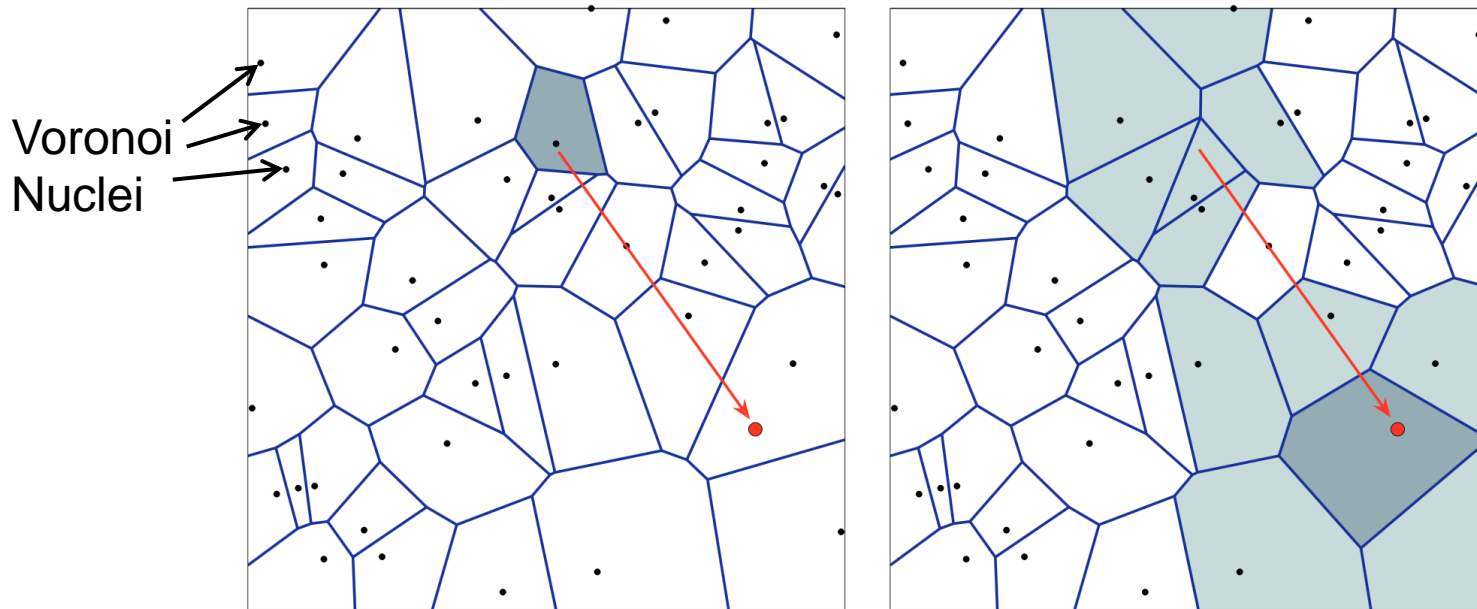
Parameterization 3 has higher likelihoods *and* uses fewer Gaussians.



Tomography

MCMC with parameterization of unknown # of Voronoi nuclei with unknown velocities and locations was proposed (Sambridge et al., 1995).

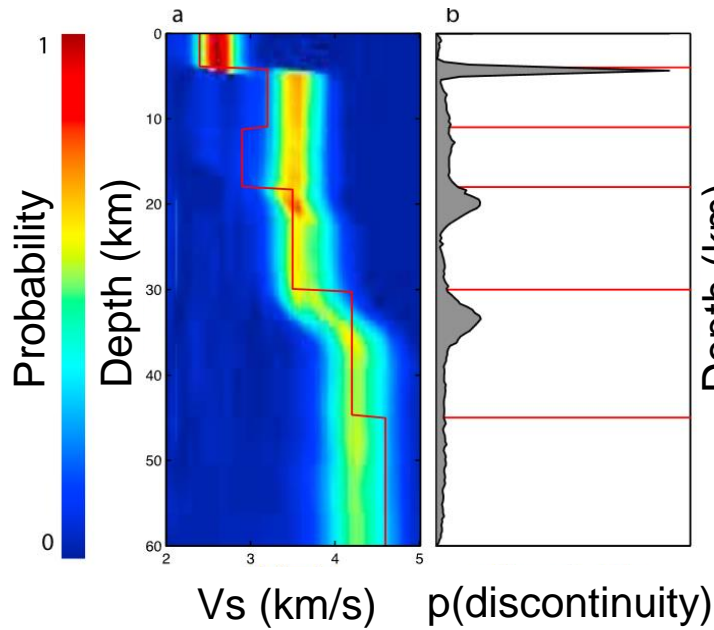
Allows for cells of varying shapes and sizes, and multiple parameterizations for the same structure.



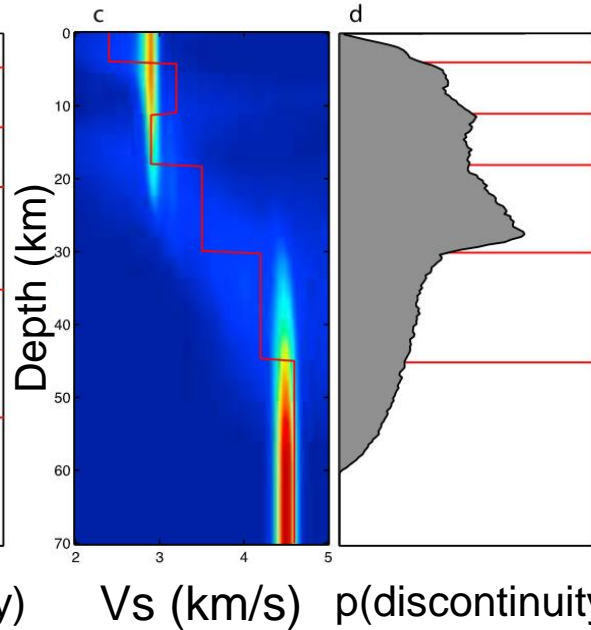
Movement of a Voronoi nucleus to a new location. The velocity structure in each of the teal cells has been changed. After Bodin et al. (2009).

Joint Inversion

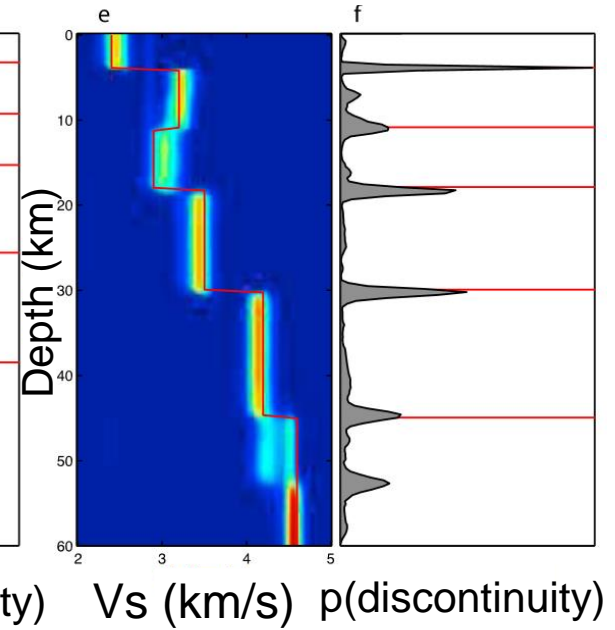
Only RF



Only SWD

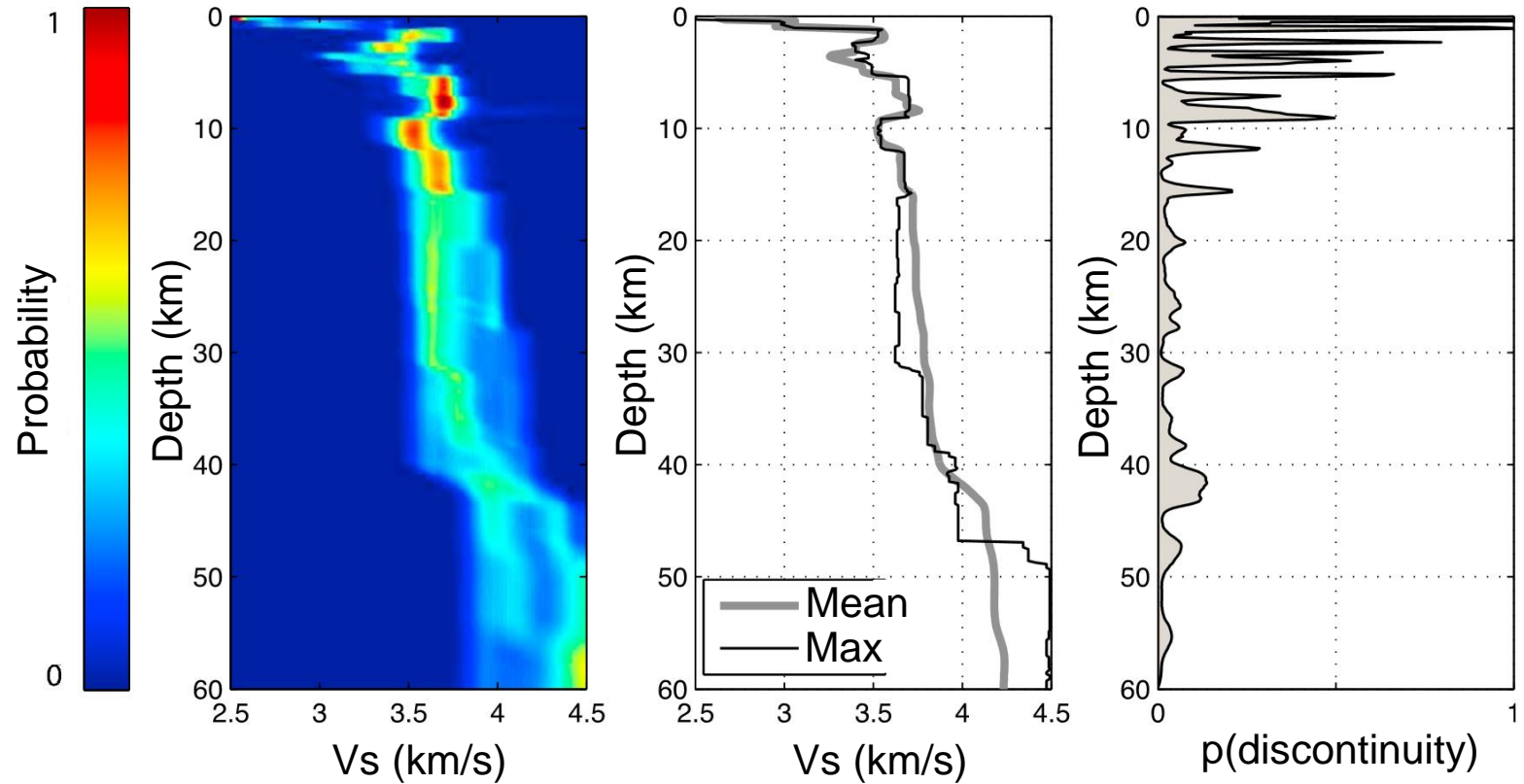


Joint (RF+SWD)



After Bodin et al., (2012)

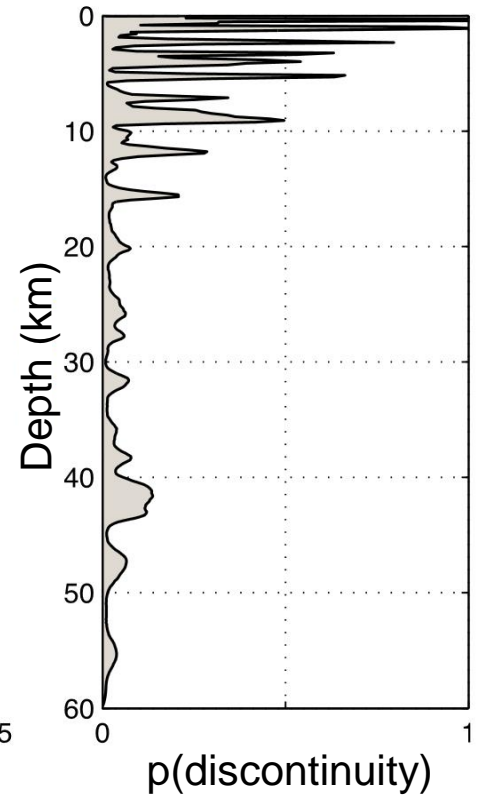
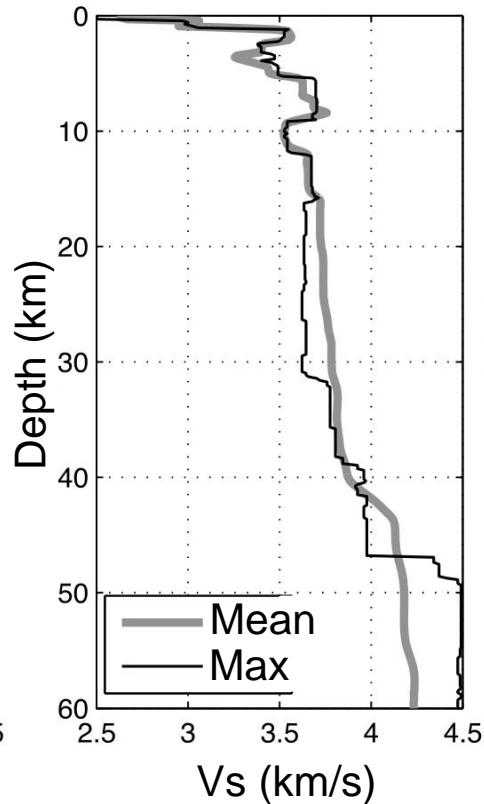
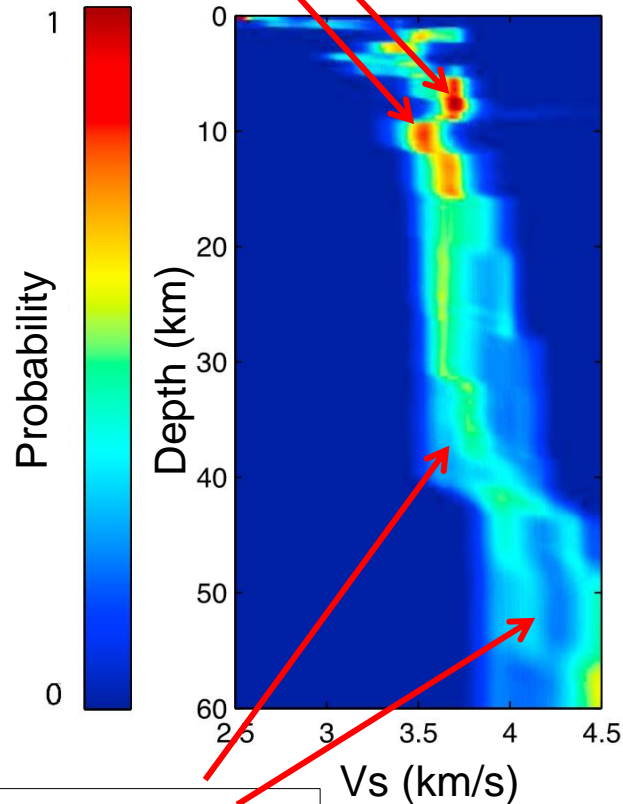
Joint Inversion



After Bodin et al., (2012)

Joint Inversion

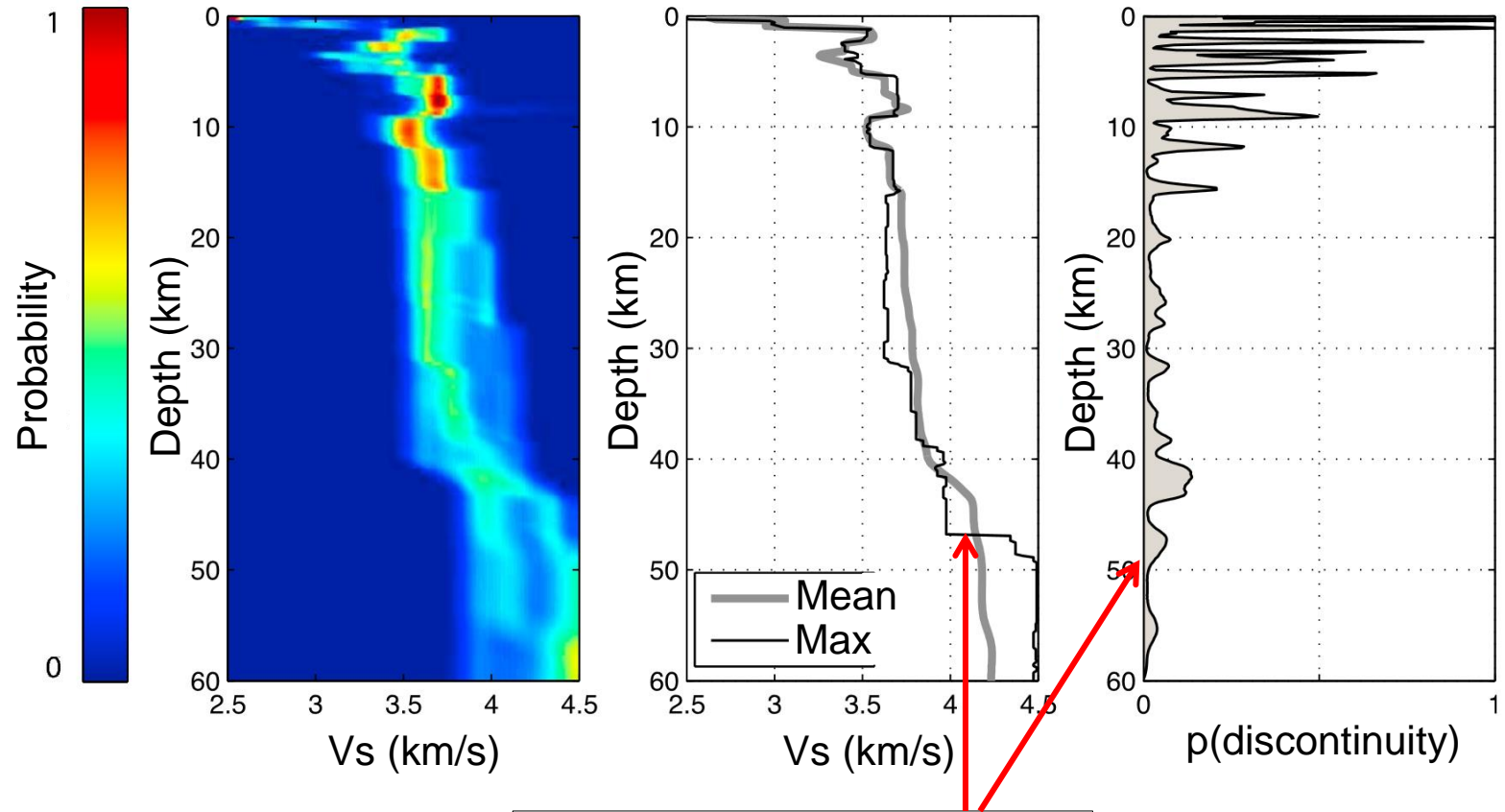
Well-Constrained Features



More uncertainty on these velocities

After Bodin et al., (2012)

Joint Inversion



Sharp increase in max. posterior model not necessarily a discontinuity

After Bodin et al., (2012)

Conclusions

Transdimensional MCMC:

- Is a data-driven approach allowing for dimensionality to be decided by the data;
- Results in an ensemble solution of models that can make uncertainty analysis simpler;
- Provides uncertainties on individual features of solutions;
- Can jump out of local minima and can sample non-unique solutions.

Acknowledgements

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Fellow “transdimensionalers”

CREWES

References

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Happy Valentine's Day!

