

Sequential Knot Selection in Sparse Gaussian Processes

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Outline

- Basics of Gaussian Processes (GPs)
- Sparse GPs using knots
- One-at-a-time (OAT) selection
- Applications

Non-parametric regression

Suppose we have the model

$$y_i = f(x_i) + \epsilon_i$$

where

- response $y_i \in \mathbb{R}$ (for simplicity)
- input $x_i \in \mathcal{X} \subset \mathbb{R}^d$
- noise $\epsilon_i \stackrel{\text{ind}}{\sim} N(0, \tau^2)$
- unknown $f : \mathcal{X} \rightarrow \mathbb{R}$

We observe pairs (y_i, x_i^\top) for $i = 1, \dots, N$ and we are interested in inference on the unknown $f(\cdot)$.

Gaussian Process

Assume a Gaussian Process (GP) prior for f :

$$f(x) \sim \mathcal{GP}(m(x), k_\theta(x, x'))$$

which assumes, for any finite subset,

$$f_x = \begin{bmatrix} f(x_1) \\ \vdots \\ f(x_M) \end{bmatrix} \sim \mathcal{N}_M(m_x, \Sigma_{xx})$$

where $m_x = [m(x_1), \dots, m(x_M)]^\top$ and

$$\Sigma_{xx}(i, j) = k_\theta(x_i, x_j)$$

for some kernel (covariance function) $k_\theta : \mathcal{X} \times \mathcal{X} \rightarrow \mathbb{R}$.

[Rasmussen and Williams, 2006]

Kernel

Kernel controls how smooth the process is both by determining:

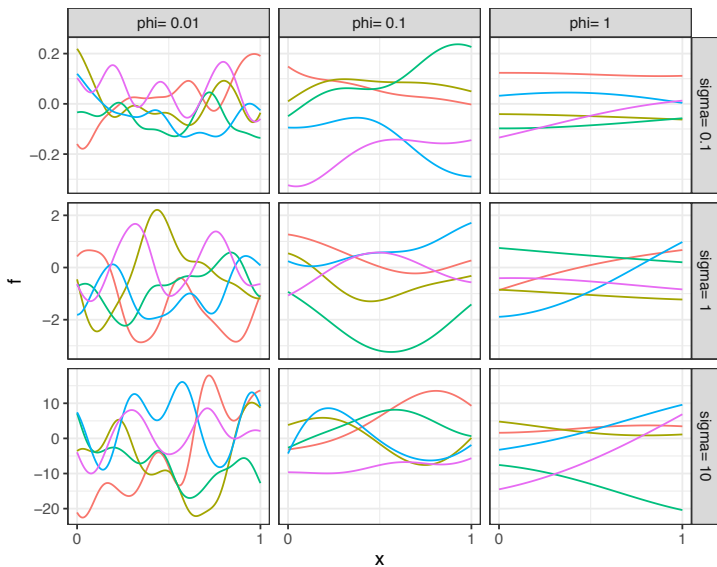
- function differentiability and
- function wiggleness.

As an example, the squared exponential (or Gaussian) kernel is

$$k_{\theta}(x_i, x_j) = \sigma^2 \exp \left(-\frac{1}{2} \frac{(x_i - x_j)^{\top} (x_i - x_j)}{\phi} \right)$$

which provides infinitely differentiable GP realizations. The parameter σ^2 is the variance that controls the overall magnitude of the function and ϕ is the length-scale that controls how wiggly the function is.

Gaussian Process Simulations (squared exponential kernel)



Training a GP

Find the maximum likelihood estimator (MLE) for $\theta = (\tau^2, \sigma^2, \phi)$,

$$\hat{\theta} = \operatorname{argmax}_{\theta} p(y|\theta) = \operatorname{argmax}_{\theta} N(y; m(x), \tau^2 \mathbf{I} + \Sigma(\theta))$$

where $y = (y_1, \dots, y_N)$. The log-likelihood is

$$\begin{aligned} \log \mathcal{N}(y; m(x), \tau^2 \mathbf{I} + \Sigma_{\theta}) = & C \\ & - \frac{1}{2} \log |\tau^2 \mathbf{I} + \Sigma(\theta)| \\ & - \frac{1}{2} (y - m_x)^{\top} [\tau^2 \mathbf{I} + \Sigma(\theta)]^{-1} (y - m_x) \end{aligned}$$

Predicting from a GP

Function estimation (prediction) from a GP is based on the following joint distribution:

$$\begin{matrix} y \\ f_{x^*} \end{matrix} \mid \hat{\theta} \sim \left(\begin{bmatrix} m_x \\ m_{x^*} \end{bmatrix}, \begin{bmatrix} \hat{\Sigma}_{xx} + \hat{\tau}^2 \mathbf{I} & \hat{\Sigma}_{xx^*} \\ \hat{\Sigma}_{x^*x} & \hat{\Sigma}_{x^*x^*} \end{bmatrix} \right)$$

where

- $x^* = (x_1^*, \dots, x_{N^*}^*)$ represents a set of prediction locations,
- $f_{x^*} = (f(x_1^*), \dots, f(x_{N^*}^*))^\top$ represents a set of prediction values,
- $m_{x^*} = (m(x_1^*), \dots, m(x_{N^*}^*))^\top$,
- $\Sigma_{x^*x^*}(i, j) = k_{\hat{\theta}}(x_i^*, x_j^*)$, and
- $\Sigma_{xx^*}(i, j) = k_{\hat{\theta}}(x_i, x_j^*)$.

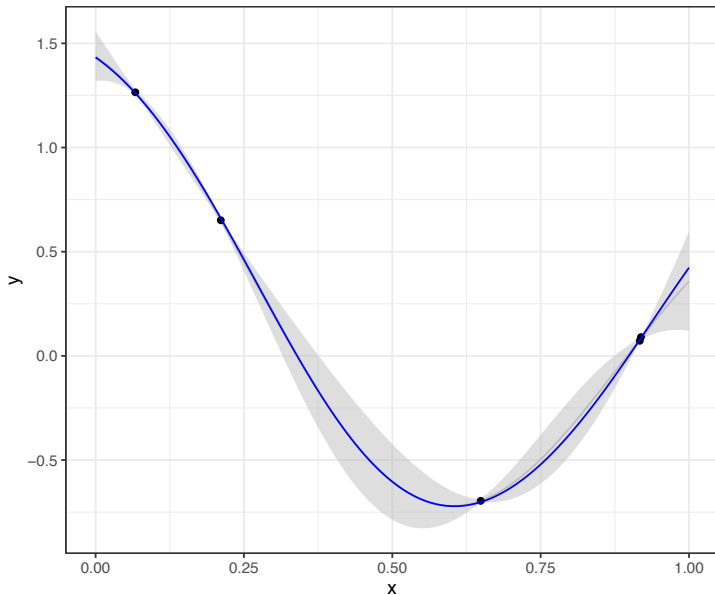
Thus, the desired conditional distribution is

$$f_{x^*} \mid y, \hat{\theta} \sim \mathcal{N}(\hat{m}_{x^*}, \hat{\Sigma}_{x^*x^*})$$

where

$$\begin{aligned} \hat{m}_{x^*} &= m_{x^*} + \hat{\Sigma}_{x^*x} \left[\hat{\tau}^2 \mathbf{I} + \hat{\Sigma}_{xx} \right]^{-1} (y - m_x) \\ \hat{\Sigma}_{x^*x^*} &= \hat{\Sigma}_{x^*x^*} - \hat{\Sigma}_{x^*x} \left[\hat{\tau}^2 \mathbf{I} + \hat{\Sigma}_{xx} \right]^{-1} \hat{\Sigma}_{xx^*}. \end{aligned}$$

Graphical representation



Training a GP - revisited

Find the maximum likelihood estimator (MLE) for $\theta = (\tau^2, \sigma^2, \phi)$,

$$\hat{\theta} = \operatorname{argmax}_{\theta} p(y|\theta) = \operatorname{argmax}_{\theta} N(y; m_x, \tau^2 \mathbf{I} + \Sigma(\theta))$$

where $y = (y_1, \dots, y_N)$. The log-likelihood is

$$\begin{aligned} \log \mathcal{N}(y; m_x, \tau^2 \mathbf{I} + \Sigma(\theta)) = & C \\ & - \frac{1}{2} \log |\tau^2 \mathbf{I} + \Sigma(\theta)| \\ & - \frac{1}{2} (y - m_x)^{\top} [\tau^2 \mathbf{I} + \Sigma(\theta)]^{-1} (y - m_x) \end{aligned}$$

If there are N observations, $\Sigma(\theta)$ is an $N \times N$ covariance matrix and thus the computational time scales as $\mathcal{O}(N^3)$.

This is doable if $N \approx 1,000$ but not when you start getting larger and larger data sets.

Fully Independent Conditional (FIC) Approximation

Introduce a set of knots $x^\dagger = \{x_1^\dagger, \dots, x_K^\dagger\}$, such that

$$p(f_x, f_{x^\dagger} | \theta) = p(f_x | f_{x^\dagger}, \theta) p(f_{x^\dagger} | \theta).$$

where

$$\begin{aligned} f_x | f_{x^\dagger}, \theta &\sim \mathcal{N}(m_x + \Sigma_{xx^\dagger} \Sigma_{x^\dagger x^\dagger}^{-1} (f_{x^\dagger} - m_{x^\dagger}), \Lambda) \\ f_{x^\dagger} | \theta &\sim \mathcal{N}(m_{x^\dagger}, \Sigma_{x^\dagger x^\dagger}) \end{aligned}$$

with $\Lambda = \text{diag}(\Sigma_{xx} - \Sigma_{xx^\dagger} \Sigma_{x^\dagger x^\dagger}^{-1} \Sigma_{x^\dagger x})$.

This joint implies the following marginal distribution for f_x :

$$f_x | \theta \sim \mathcal{N}(m_x, \Lambda + \Sigma_{xx^\dagger} \Sigma_{x^\dagger x^\dagger}^{-1} \Sigma_{x^\dagger x})$$

which has the correct marginal means and variances, but the covariances are controlled by the knots.

[Seeger et al., 2003, Quiñero-Candela and Rasmussen, 2005, Snelson and Ghahramani, 2006, Banerjee et al., 2008, Finley et al., 2009, Titsias, 2009, Cao et al., 2013]

Train FIC Model

Let $\Psi_{xx} \equiv \Lambda(\theta) + \Sigma_{xx^\dagger}(\theta)\Sigma_{x^\dagger x^\dagger}(\theta)^{-1}\Sigma_{x^\dagger x}(\theta)$, then

$$Y|x^\dagger, \theta \sim \mathcal{N}(m_x, \tau^2 I + \Psi_{xx}).$$

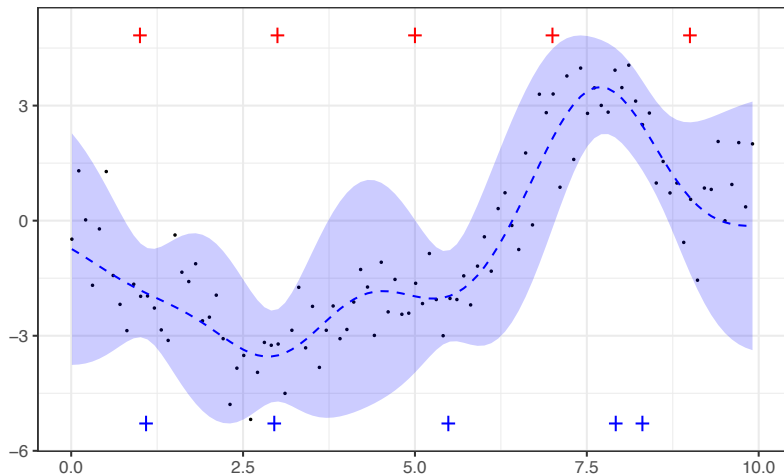
Train the model by finding

$$\hat{x}^\dagger, \hat{\theta} = \operatorname{argmax}_{x^\dagger, \theta} \mathcal{N}(y; m_x, \tau^2 I + \Psi_{xx}).$$

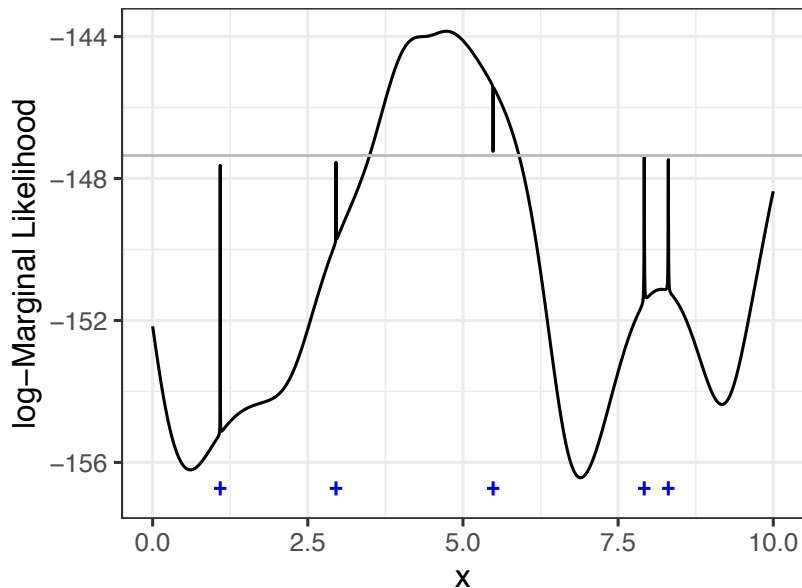
Appealing due to similarity with Full GP MLE approach, but there are a number of questions:

- how many knots are needed?
- where should we initialize the knots?
- when do we stop our iterative optimization algorithm?

Simultaneous knot optimization



Adding another knot



Knot selection algorithm

Algorithm 1. OAT knot selection algorithm. Convergence in the repeat loop is declared when the change in the objective function, the log-marginal likelihood, falls below a threshold. Set initial number of knots (K_I).

```

1 Initialize:  $x^\dagger = \{x_i^\dagger\}_{i=1}^{K_I}$  ;
2  $\hat{\theta} = \operatorname{argmax}_{\theta} p(y|x, x^\dagger, \theta)$  ;
3 repeat
4   | propose new knot  $x^{\dagger*} \leftarrow J(y, x, x^\dagger, \hat{\theta})$  ;
5   |  $(\hat{x}^{\dagger*}, \hat{\theta}) = \operatorname{argmax}_{(x^{\dagger*}, \theta)} p(y|x, \{x^\dagger, x^{\dagger*}\}, \theta)$  ;
6   |  $x^\dagger = \{x^\dagger, \hat{x}^{\dagger*}\}$  ;
7 until  $|x^\dagger| = K_{max}$  or convergence;
```

Bayesian optimization

Let

- $w_{1:t-1}$ be the vector of log-marginal likelihood values at the candidates for the knot proposal which have thus far been explored at time t
- $w^+ = \max(w_{1:t-1})$

Let $W(z)$ be the unknown marginal likelihood at input location z , then expected improvement is

$$\alpha\left(z; w_{1:t-1}, \left\{x_1^\dagger, \dots, x_{t-1}^\dagger\right\}\right) = (E[W(z)|w_{1:t-1}] - w^+) \Phi\left(\frac{E[W(z)|w_{1:t-1}] - w^+}{\sqrt{V[W(z)|w_{1:t-1}]}}\right) + \sqrt{V[W(z)|w_{1:t-1}]} \phi\left(\frac{E[W(z)|w_{1:t-1}] - w^+}{\sqrt{V[W(z)|w_{1:t-1}]}}\right).$$

where ϕ and Φ are the pdf and cdf of a standard normal, respectively.

We will model the unknown marginal likelihood $W(z)$ using a **meta GP**.

Knot proposal algorithm

Algorithm 2. Knot proposal algorithm. Set the minimum (T_{min}) and maximum (T_{max}) number of marginal likelihood evaluations.

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-
- 1 set the mean of the meta GP equal to $\log p\left(y \mid x, \{x^\dagger, \cdot\}, \hat{\theta}\right)$;
 - 2 sample $x_1^\dagger, \dots, x_{T_{min}}^\dagger$ without replacement from x ;
 - 3 augment known marginal likelihood values $w_j = \log p\left(y \mid x, \{x^\dagger, x_j^\dagger\}, \hat{\theta}\right)$ for $j = 1, \dots, k$ with evaluations of the marginal likelihood at the new knots, that is $w_{k+j} = \log p\left(y \mid x, \{x^\dagger, x_j^\dagger\}, \hat{\theta}\right)$ for $j = 1, \dots, T_{min}$;
 - 4 **for** $t = T_{min} + 1, \dots, T_{max}$ **do**
 - 5 update covariance parameters in meta GP ;
 - 6 $x_t^* = \operatorname{argmax}_{z \in x \setminus \{x_l^\dagger\}_{l=1}^{t-1}} \alpha\left(z; w, \{x_1^\dagger, \dots, x_{t-1}^\dagger\}\right)$;
 - 7 $w_t = \log p\left(y \mid x, \{x^\dagger, x_t^*\}, \hat{\theta}\right)$;
 - 8 **end**
 - 9 return x_j^* such that $j = \operatorname{argmax}_t w_t$
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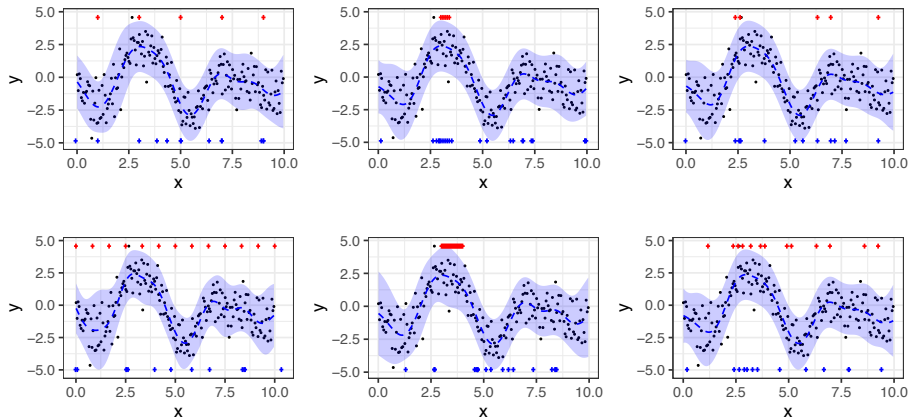
Knot selection algorithm

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```

1 Initialize:  $x^\dagger = \{x_i^\dagger\}_{i=1}^{K_I}$  ;
2  $\hat{\theta} = \operatorname{argmax}_{\theta} p(y|x, x^\dagger, \theta)$  ;
3 repeat
4   | propose new knot  $x^{\dagger*} \leftarrow J(y, x, x^\dagger, \hat{\theta})$  ;
5   |  $(\hat{x}^{\dagger*}, \hat{\theta}) = \operatorname{argmax}_{(x^{\dagger*}, \theta)} p(y|x, \{x^\dagger, x^{\dagger*}\}, \theta)$  ;
6   |  $x^\dagger = \{x^\dagger, \hat{x}^{\dagger*}\}$  ;
7 until  $|x^\dagger| = K_{max}$  or convergence;
```

One-D Gaussian data



Starting locations: evenly spaced (left), adversarial (middle), random (right)
 Algorithm: OAT (top) and simultaneous (bottom)

Computational results

Method	Initialization	K	RMSE	Runtime	GA Steps	log-Likelihood
Full GP	–	–	0.192	–	–	-311.720
OAT	Uniform	13	0.180	50	464	-308.120
OAT	Adversarial	22	0.228	96	669	-308.587
OAT	Random	13	0.228	50	470	-308.225
Simult.	Uniform	13	0.220	140	212	-306.852
Simult.	Adversarial	22	0.196	700	529	-308.398
Simult.	Random	13	0.247	88	140	-308.071

Performance metrics

All data models:

$$MNLP = \text{median}_{i \in 1, \dots, N_{test}} \{-\log p(\tilde{y}_i | x^\dagger, \hat{\theta}, y)\}.$$

$$AUKL = \frac{1}{N_{test}} \sum_{i=1}^{N_{test}} \int p_{full}(f(\tilde{x}_i) | \hat{\theta}, y) \log \frac{p_{full}(f(\tilde{x}_i) | \hat{\theta}, y)}{p_{sparse}(f(\tilde{x}_i) | x^\dagger, \hat{\theta}, y)} df(\tilde{x}_i).$$

Gaussian:

$$SRMSE = \sigma_{\tilde{y}}^{-1} \sqrt{\frac{1}{N_{test}} \sum_{i=1}^{N_{test}} (E[f(\tilde{x}_i) | Y] - \tilde{y}_i)^2},$$

where $\sigma_{\tilde{y}}^2 = \frac{1}{N_{test}-1} \sum_{i=1}^{N_{test}} (\tilde{y}_i - \underline{\tilde{y}})^2$, $\underline{\tilde{y}} = \frac{1}{N_{test}} \sum_{i=1}^{N_{test}} \tilde{y}_i$, and \tilde{y} is the vector of test set target values.

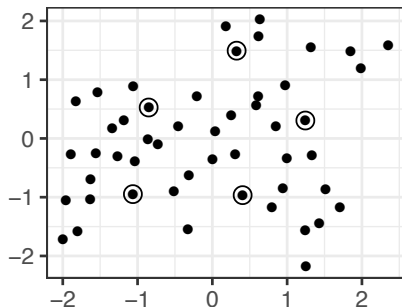
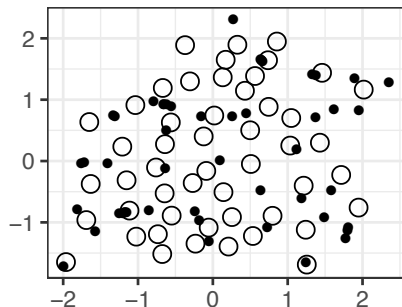
Boston Housing

490 observations (random 80% for training) with $d = 3$

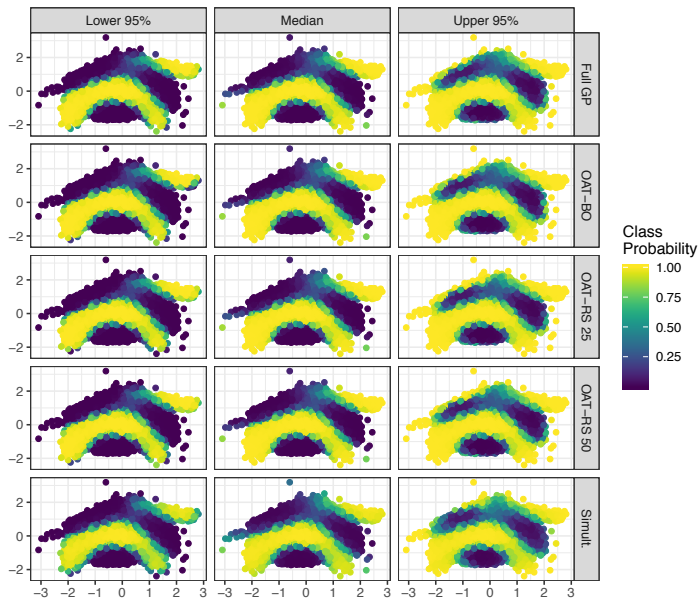
Method	Runtime	K	K/Tmax	SRMSE	MNLP	AUKL
Full	394	–	–	0.359	2.500	0.000
OAT-BO	545	13	25	0.366	2.466	0.045
OAT-RS	356	12	25	0.366	2.464	0.039
OAT-RS	339	15	50	0.364	2.469	0.047
Simult.	25831	50	–	0.378	2.291	0.356
Simult.	3945	13	–	0.356	2.313	0.242

Banana data

Binary classification: 5300 observations (random 10% as training) with $d = 3$



Locations of initialized and estimated knots for the simultaneously optimized model with 50 knots (left) and for the OAT-BO model (right). Open circles are initial knots and solid points are estimated knots.



Method	Runtime	Tmax	K	MNLP	AUKL
Full	26795	–	–	0.038	0.000
OAT-BO	3150	25	50	0.037	0.061
OAT-RS	2954	25	50	0.038	0.051
OAT-RS	3471	50	50	0.038	0.039
Simult.	6219	–	50	0.069	3.265

Summary

One-at-a-time (OAT) knot selection

- Similar predictive performance to simultaneous knot selection
- Better represents full GP compared to simultaneous knot selection
- Better computational efficiency than simultaneous knot selection

These slides are available

- <https://github.com/jarad/SFU2020>
- <http://www.jarad.me/research/presentations.html>

Thank you!

Other links:

- <http://www.jaradniemi.com/>
- <https://www.youtube.com/jaradniemi>
- <https://twitter.com/jaradniemi>

Reference

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