Kalman Filter

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Outline

- Introduction
- 2 Details
 - Step 1. Time dynamic systems
 - Step 2. Bayes Theorem
 - Step 3. Recursion
 - Step 4. Solutions
- 3 Conclusion

Overview

- The Kalman filter is a <u>solution</u> to the <u>recursive</u>
 Bayesian equations of a linear time-dynamic system
- Ok, so working backwards, we need to know about time dynamic systems, Bayesian equations, recursion and a solution to these equations:
 - Time dynamic systems
 - 2 Bayes Theorem (Prior, Likelihood, Posterior)
 - Recursion (Predict and Update)
 - Solutions (HMM, Kalman Filter, PDAF, JPDAF, Particle Filter)
- Let's look at these one at a time



Step 1. Time dynamic systems Step 2. Bayes Theorem

tep 3. Recursion

Time dynamic systems

• Time-dynamic systems are described by two equations:

$$\mathbf{x}_k = f_k(\mathbf{x}_{k-1}, \mathbf{v}_{k-1})$$

 $\mathbf{z}_k = h_k(\mathbf{x}_k, \mathbf{n}_k)$

- States are denoted by x
- Observations are denoted by z
- \bullet Process noise is denoted by ${\bf v}$
- Observation noise is denoted by n
- Time is denoted by k
- State prediction model, $f_k: R^D \times R^D \to R^D$, describes state evolution
- Observation model, $h_k : R^N \times R^N \to R^N$, relates observations to the states



Step 1. Time dynamic systems

Step 3. Recursio

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Time dynamic systems Examples

- Robotics: motion of a robot
- Agriculture: color change of a fruit
- Human body: Secretion of insulin
- Finance: Fluctuations in inflation
- Mechanical: Spring mass system
- Radars: Motion of an aircraft

Step 1. Time dynamic system Step 2. Bayes Theorem

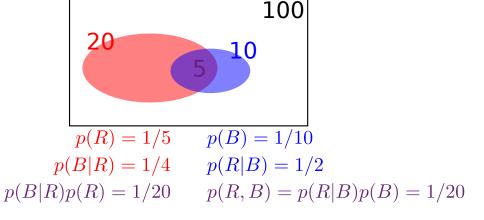
Step 4. Solutions

Bayes Theorem

Ok, time dynamic systems make sense but how does the Bayes theorem fit into our picture here?

Step 2. Bayes Theorem

Bayes Theorem Graphical demonstration



Step 1. Time dynamic system Step 2. Bayes Theorem

Step 4. Solutions

Bayes Theorem

Example from real life: Prosecutor's fallacy

- If someone is guilty, the probability of a DNA match is large, p(D|g) is large
 - convict?
- Probability that a person is guilty if picked randomly from a large database is small, p(g) is very small
- Probability p(D,g) = p(D|g)p(g) is not so large after all
- So the prior information, p(g) is important!

Step 1. Time dynamic systems
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Bayes Theorem

Relationship with time dynamic systems

Posterior distribution of states (our goal)
Prior distribution of states (our belief)
Likelihood distribution (we can compute it)
Observation probability (we won't need it!)

• Now, what's the deal with recursion?

ep 1. Time dynamic systems

Step 3. Recursion

Recursion

- In Bayes' theorem, we talked about 3 important probability distributions (posterior, prior, likelihood)
- We want to compute the posterior distribution at every time instant
- This is recursion, incremental updates, not batch mode

ep 1. Time dynamic system:

Step 3. Recursion

Recursion

- We assume that our system and observation models are available in probabilistic form
- The probabilistic state-space formulation and the requirement for updating of information on receipt of new measurements are ideally suited for the Bayesian approach
 - Prediction: predict states using model
 - Update: correction applied to prediction after observation arrives

tep 1. Time dynamic systems

Step 3. Recursion

Recursion

a. Predict

x _k	state at time <i>k</i>
Z_{k-1}	all observations till time $k-1$

$$\begin{split} \frac{p(x_k|Z_{k-1})}{p(x_k|Z_{k-1})} &= \frac{p(x_k,Z_{k-1})}{p(Z_{k-1})} \\ &= \frac{\int p(x_k,x_{k-1},Z_{k-1})dx_{k-1}}{p(Z_{k-1})} \\ &= \frac{\int p(x_k|x_{k-1},Z_{k-1})p(x_{k-1},Z_{k-1})dx_{k-1}}{p(Z_{k-1})} \\ &= \frac{\int p(x_k|x_{k-1},Z_{k-1})p(x_{k-1},Z_{k-1})dx_{k-1}}{p(Z_{k-1})} \\ &= \frac{\int p(x_k|x_{k-1})p(x_{k-1}|Z_{k-1})p(Z_{k-1})dx_{k-1}}{p(Z_{k-1})} \\ &= \int p(x_k|x_{k-1})p(x_{k-1}|Z_{k-1})dx_{k-1} \end{split}$$

- Step 3. Recursion

Recursion b. Update

$$\begin{split} p(x_k|Z_k) &= \frac{p(x_k,Z_k)}{p(Z_k)} \\ p(x_k|Z_k) &= \frac{p(x_k,Z_k,Z_{k-1})}{p(z_k,Z_{k-1})} \\ p(z_k,Z_{k-1}) &= \frac{p(z_k|x_k,Z_{k-1})}{p(z_k,Z_{k-1})} \\ &= \frac{p(z_k|x_k,Z_{k-1})}{p(z_k|Z_{k-1})p(Z_{k-1})} \\ &= \frac{p(z_k|x_k)p(x_k|Z_{k-1})p(Z_{k-1})}{p(z_k|Z_{k-1})p(Z_{k-1})} \\ &= \frac{p(z_k|x_k)p(x_k|Z_{k-1})p(Z_{k-1})}{p(z_k|x_k)p(x_k|Z_{k-1})} \\ &= \frac{p(z_k|x_k)p(x_k|Z_{k-1})}{p(z_k|x_k)p(x_k|Z_{k-1})} \\ &= \frac{p(z_k|x_k)p(x_k|Z_{k-1})}{p(z_k|x_k)p(x_k|Z_{k-1})} \end{split}$$

ep 1. Time dynamic system: ep 2. Baves Theorem

Step 3. Recursion Step 4. Solutions

Solutions

How to implement?

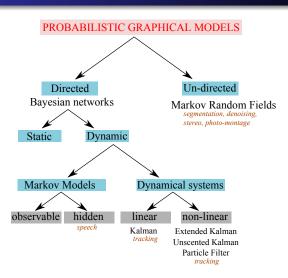
- The recursive propagation of the state density is only a conceptual solution in that in general, it cannot be determined analytically.
- Solutions do exist in a restrictive set of cases, including the Kalman filter and grid-based filter
 - State space is discrete and consists of finite number of states
- In many situations of interest, the conditions of linearity and Gaussianity do not hold, and approximations are necessary.
- Three approximate nonlinear Bayesian filters are the Extended Kalman Filter (EKF), approximate grid based methods and the particle filter

Step 2 Rayes Theorem

Step 3 Recursion

Step 3. Recursion Step 4. Solutions

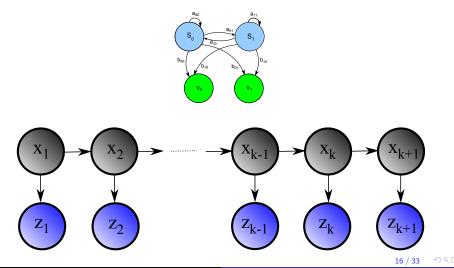
Solutions Bigger picture



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Solutions

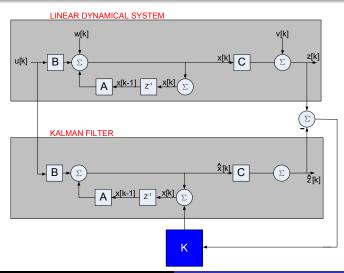
a. Hidden Markov Models



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Solutions

b. Kalman Filter



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b. Kalman Filter

- Prior
 - Past information through time k-1
 - summarized approximately by a sufficient statistic in the form of a Gaussian posterior

$$p(x_{k-1|k-1}|Z_{k-1}) = \mathcal{N}(\hat{x}_{k-1|k-1}, P_{k-1|k-1})$$

- Prediction
 - The state prediction is distributed as,

$$p(x_{k|k-1}|Z_{k-1}) = \mathcal{N}(\hat{x}_{k|k-1}, P_{k|k-1})$$

The observation prediction is distributed as,

$$p(z_{k|k-1}) = \mathcal{N}(\hat{z}_{k|k-1}, \mathcal{S}_k)$$

- Update
 - innovation
 - Kalman gain
 - aposteriori state estimate and aposteriori covariance estimate



Step 4. Solutions

Solutions

b. Kalman Filter

PREDICTION

1a. state
$$\hat{\mathbf{x}}_{k|k-1} = \mathbf{A}_k \hat{\mathbf{x}}_{k-1} + \mathbf{B}_k \mathbf{u}_k$$

1b. state covariance
$$\mathbf{M}_{k|k-1} = \mathbf{A}_k \mathbf{M}_{k-1|k-1} \mathbf{A}_k^T + \mathbf{Q}_k$$

2a. measurement
$$\hat{\mathbf{z}}_{k|k-1} = \mathbf{C}_k \hat{\mathbf{x}}_{k|k-1}$$

2b. measurement covariance (called innovation covariance)
$$\mathbf{S}_k = \mathbf{C}_k \mathbf{M}_{k|k-1} \mathbf{C}_k^T + \mathbf{R}_k$$

3. gain
$$\mathbf{K}_k = \mathbf{M}_{k|k-1} \mathbf{C}_k^T \mathbf{S}_k^{-1}$$

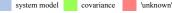
observation is received

4. innovation
$$\mathbf{y}_k = \mathbf{z}_k - \hat{\mathbf{z}}_{k|k-1}$$

UPDATE

5a. state
$$\hat{\mathbf{x}}_{k|k-1} = \hat{\mathbf{x}}_{k|k-1} + \mathbf{K}_k \mathbf{y}_k$$

5b. state covariance
$$\mathbf{M}_{k|k-1} = (\mathbf{I} - \mathbf{K}_k \mathbf{C}_k) \mathbf{M}_{k|k-1}$$



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c. PDAF: Extension of Kalman Filter

Probabilistic Data Association Filter

- Computationally efficient
- Data association
- Single targets in clutter, high false alarm rate
- Extension of Kalman filter
 - Primary difference: computation of innovations
 - Calculates association probabilities to the target for each validated measurement

Step 3. Recursion

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c. PDAF: Extension of Kalman Filter

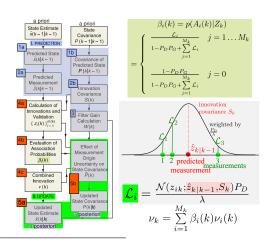
Assumptions:

- Number of targets
 - Single target tracking
- Palse alarms
 - At most, one of the validated measurements can be target originated
 - Remaining measurements
 - incorrect, i.e., false alarms
 - i.i.d
 - \bullet parametric form: Poisson distribution with known spatial density λ
 - non-parametric form: uniform spatial distribution

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c. PDAF: Extension of Kalman Filter



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c. PDAF: Extension of Kalman Filter

PDAF + multiple model Kalman Filters :



US Navy ROHR (Relocatable Over the Horizon Radar) for long-range surveillance for drug interdiction against aircraft and ships

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d IPDAF: Extension of Kalman Filter

Joint Probabilistic Data Association Filter

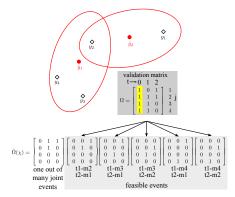
- Extension of PDAF to multi-target tracking
 - ullet Only difference: computation of association probabilities, eta_{ik}
- Handles multiple targets by considering all measurements for all targets.
- Probability density of each candidate measurement
 - Based on all close-by targets

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d. JPDAF: Extension of Kalman Filter

Association probabilities



$$\begin{aligned} p[\chi|Z_k] &= \frac{1}{c} \prod_{j:\tau_j=1} \mathbb{N}[z_j(k); \hat{z}^{t_j}(k|k-1), S^{t_j}(k)] \prod_{t:\delta_t=1} P_D^t \prod_{t:\delta_t=0} (1 - P_D^t) \\ \beta_j^t &= \sum_t p[\chi|Z_k] \hat{\omega}_{jt}(\chi) \end{aligned}$$

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d. JPDAF: Extension of Kalman Filter

Nearest neighbor JPDAF + EKF:



THAAD, Theater High Altitude Area Defense)



Cobra Dane, long-range surveillance against ICBMs



SBX, long-range surveillance against ICBMs

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e. Multi-hypothesis tracker

- Clustering
 - A new measurement is associated with a cluster if it falls within the validation region of any target within the cluster
- Hypothesis Generation
 - New hypotheses are generated for the measurements associated with each cluster
 - The probability of each hypothesis is calculated
 - Target estimates are then updated
- Reduction
 - In this step, hypotheses are combined or eliminated
- Simplify hypothesis matrix
 - Targets that are uniquely associated are removed from the hypothesis matrix



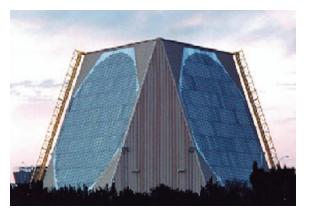
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e. Multi-hypothesis tracker

MHT + EKF:



UEWR, Upgraded Early Warning Radar, long-range surveillance against ICBMs

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Solutions f. Particle filter

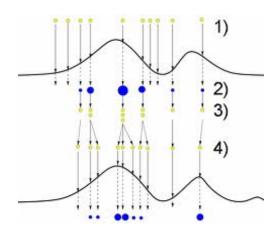
- All densities are modeled using samples
- Two step process, predict, and then update using resampling

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Solutions f. Particle filter



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Solutions f. Particle filter

- There is an essential duality between a sample and the density (distribution) from which it is generated
 - The density generates the sample
 - Conversely, given a sample, we can approximately recreate the density
- In terms of densities, the inference process is encapsulated in the updating of the prior density p(x) to the posterior density p(x|z) through the medium of the likelihood function p(z|x)
- In terms of samples, this corresponds to the updating of a sample from p(x) to a sample from p(x|z) through the likelihood function p(z|x)





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Solutions f. Particle filter

- In many application areas, elements of nonlinearity and non-Gaussianity need to be included to accurately model the underlying dynamics of a physical system
- Moreover, it is typically crucial to process data on-line as it arrives
- Particle filters are Sequential Monte Carlo (SMC) methods based on point-mass (or "particle") representations of probability densities which can be applied to any state-space model
- The particle filter generalizes the Kalman filter
- The algorithm used is Sequential Importance Sampling (SIS)



Conclusion

- There are many solutions to the Bayesian recursion equations
- The Kalman Filter is optimal if these equations are linear and the models are Gaussian

