

Robotics 2

Data Association

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Data Association

“Data association is the process of associating uncertain measurements to known tracks.”

- **Problem types**

- Track creation, maintenance, and deletion
- Single or multiple sensors
- Target detection
- False alarm model and rates
- Single or multiple targets

- **Approaches**

- **Bayesian:** compute a full (or approx.) distribution in DA space from priors, posterior beliefs, and observations
- **Non-Bayesian:** compute a maximum likelihood estimate from the possible set of DA solutions

Data Association

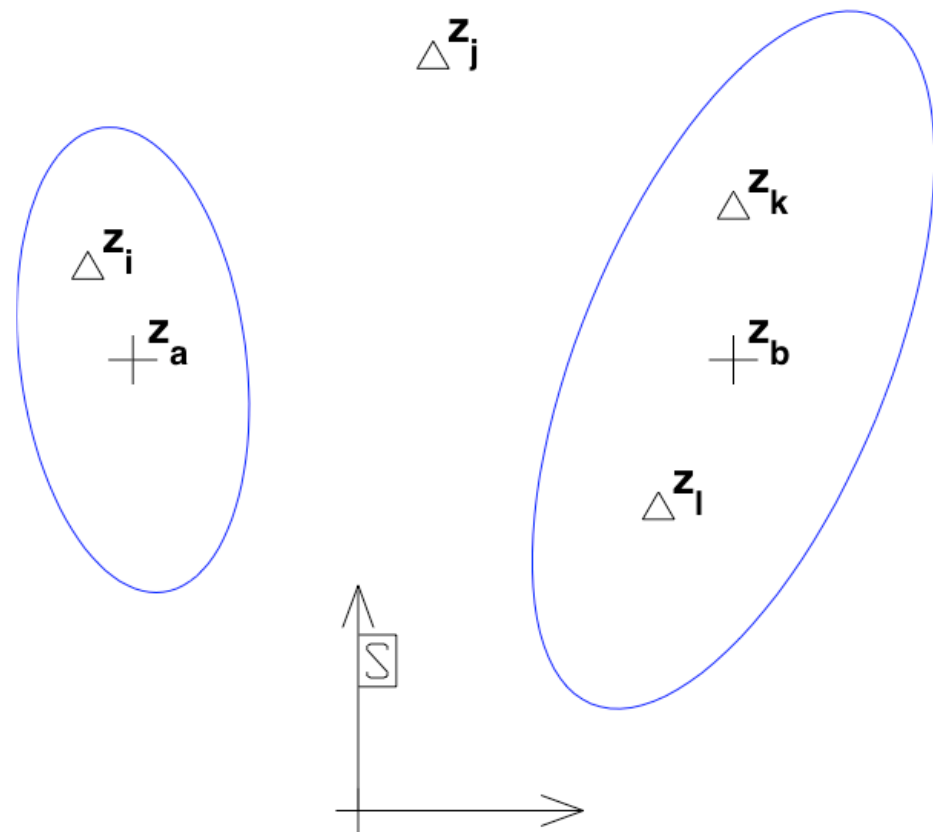
Overall procedure:

- **Make observations** (= measurements).
Measurements can be raw data (e.g. processed radar signals) or the output of some target detector (e.g. people detector)
- **Predict the measurements** from the predicted tracks.
This yields an area in sensor space where to expect an observation. The area is called **validation gate** and is used to narrow the search
- **Check if a measurement lies in the gate.**
If yes, then it is a valid candidate for a pairing/match

Data Association

What makes this a difficult problem

- **Multiple targets**
- **False alarms**
- **Detection uncertainty**
(occlusions, sensor failures, ...)
- **Ambiguities**
(several measurements in the gate)



Measurement Prediction

- Measurement and measurement cov. prediction
 - This is typically a frame transformation into sensor space

$$\hat{z}(k) = H(k)\hat{x}(k|k-1)$$

$$\hat{R}(k) = H(k)\hat{P}(k|k-1)H^T(k)$$

- If only the **position** of the target is observed (typical case), the measurement matrix is

$$\mathbf{z} = \begin{bmatrix} x & y \end{bmatrix}^T \quad H = \begin{bmatrix} 1 & 0 & 0 & \cdots \\ 0 & 1 & 0 & \cdots \end{bmatrix}$$

- Note: One can also observe
 - Velocity (Doppler radar)
 - Acceleration (accelerometers)

Validation Gate

- Assume that measurements are distributed according to a Gaussian, centered at the measurement prediction $\hat{z}(k)$ with covariance $\hat{S}(k)$

$$p(z(k)) = \mathcal{N}(z(k); \hat{z}(k), \hat{S}(k))$$

This is the **measurement likelihood model**

- Let further

$$d = \sqrt{(\mathbf{x} - \mu)^T \mathbf{C}^{-1} (\mathbf{x} - \mu)}$$

be the **Mahalanobis distance** between \mathbf{x} and μ

Validation Gate

- Then, the measurements will be in the area

$$\begin{aligned}\mathcal{V}(k, \gamma) &= \{z : (z - \hat{z})^T \hat{S}^{-1} (z - \hat{z}) \leq \gamma\} \\ &= \{z : d^2 \leq \gamma\}\end{aligned}$$

with a probability defined by the gate threshold γ
(omitting indices k)

- This area is called **validation gate**
- The threshold is obtained from the inverse χ^2 cumulative distribution at a **significance level** α
- χ^2 = “chi square”

Validation Gate

- The **shape** of the validation gate is a hyper-ellipsoid (an ellipse in 2d)
- This follows from setting

$$c = \frac{1}{(2\pi)^{k/2} |S|^{1/2}} \exp \left(-\frac{1}{2} (z - \hat{z})^T S^{-1} (z - \hat{z}) \right)$$

which gives

$$c' = (z - \hat{z})^T S^{-1} (z - \hat{z})$$

- The gate is defined by an **iso-probability contour** obtained when intersecting a Gaussian with a hyper-plane.

Validation Gate

Why a χ^2 distribution?

- Let X_i be a set of k i.i.d. standard normally distributed random variables, $X_i \sim \mathcal{N}(x; 0, 1)$.

Then, the variable Q

$$Q = \sum_{i=1}^k X_i^2$$

follows a χ^2 distribution with k “degrees of freedom”

- We will now show that the Mahalanobis distance is a sum of squared standard normally distributed RVs.

Validation Gate in 1D

- Assume 1D measurements and $\mu = \hat{z}(k)$, $\sigma^2 = \hat{S}(k)$
- The Mahalanobis distance is then

$$d^2 = (z - \mu)^T (\sigma^2)^{-1} (z - \mu) = \frac{(z - \mu)^2}{\sigma^2}$$

- By changing variables, $y = (z - \mu)/\sigma$, we have

$$y \sim \mathcal{N}(0, 1)$$

- Thus, $d^2 = y^2$ and is χ^2 distributed with 1 degree of freedom

Validation Gate in ND

- Assume ND measurements and $\mu = \hat{z}(k)$, $\Sigma = \hat{S}(k)$
- The Mahalanobis distance is then

$$d^2 = (z - \mu)^T \Sigma^{-1} (z - \mu)$$

- By changing variables, $y = C^{-1}(z - \mu)$, $\Sigma = CC^T$ we have $y \sim \mathcal{N}(0, I)$ and therefore

$$d^2 = y^T I^{-1} y \quad \Rightarrow \quad d^2 = \sum_{i=1}^k y_i^2$$

which is χ^2 distributed with k degrees of freedom.
(C is obtained from a Cholesky decomposition)

Validation Gate

Where does the threshold γ come from?

- γ , often denoted $\chi^2_{k,\alpha}$, is taken from the inverse χ^2 cumulative distribution at a level α and k d.o.f.s
- The values are typically given in tables, e.g. in most statistics books (or by the Matlab function `chi2inv`)
- Given the level α , we can now understand the interpretation of the validation gate:

The validation gate is a **region of acceptance** such that $100(1 - \alpha)\%$ of **true measurements** are **rejected**

- Typical values for α are 0.95 or 0.99

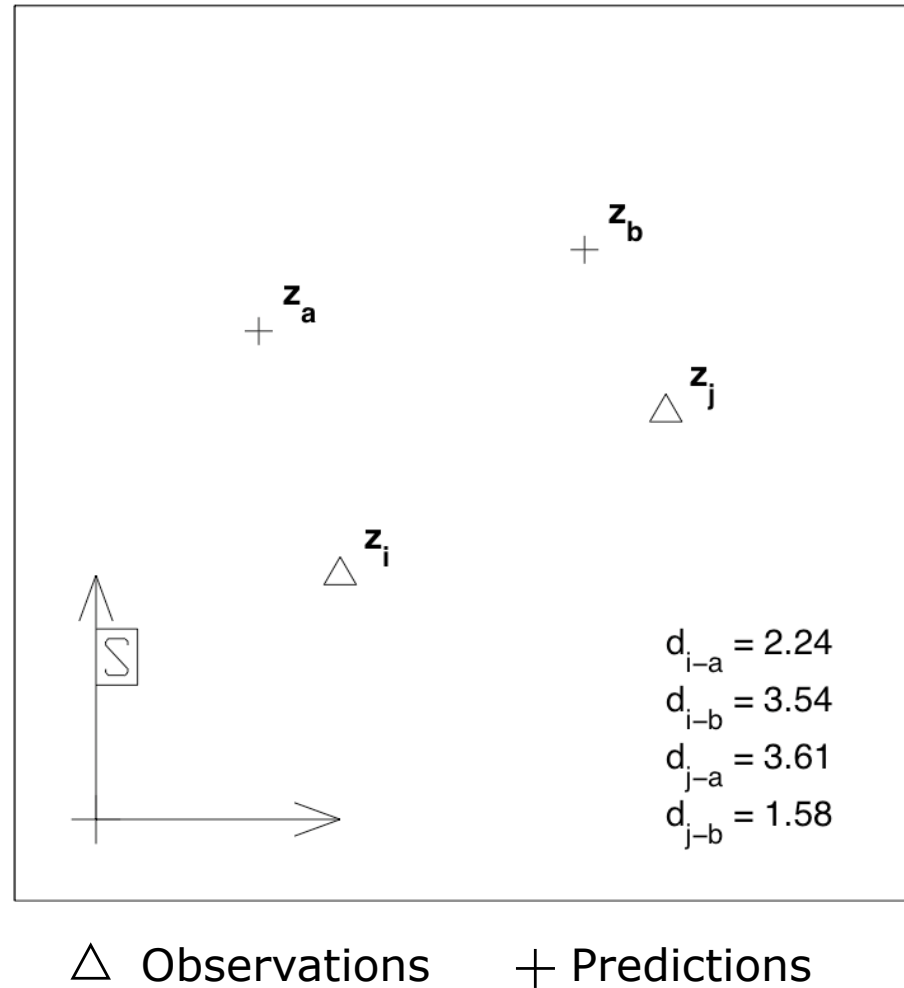
Validation Gate

Euclidian distance

Takes into account:

- ✓ Position
- ✗ Uncertainty
- ✗ Correlations

→ It seems that i-a and j-b belong together



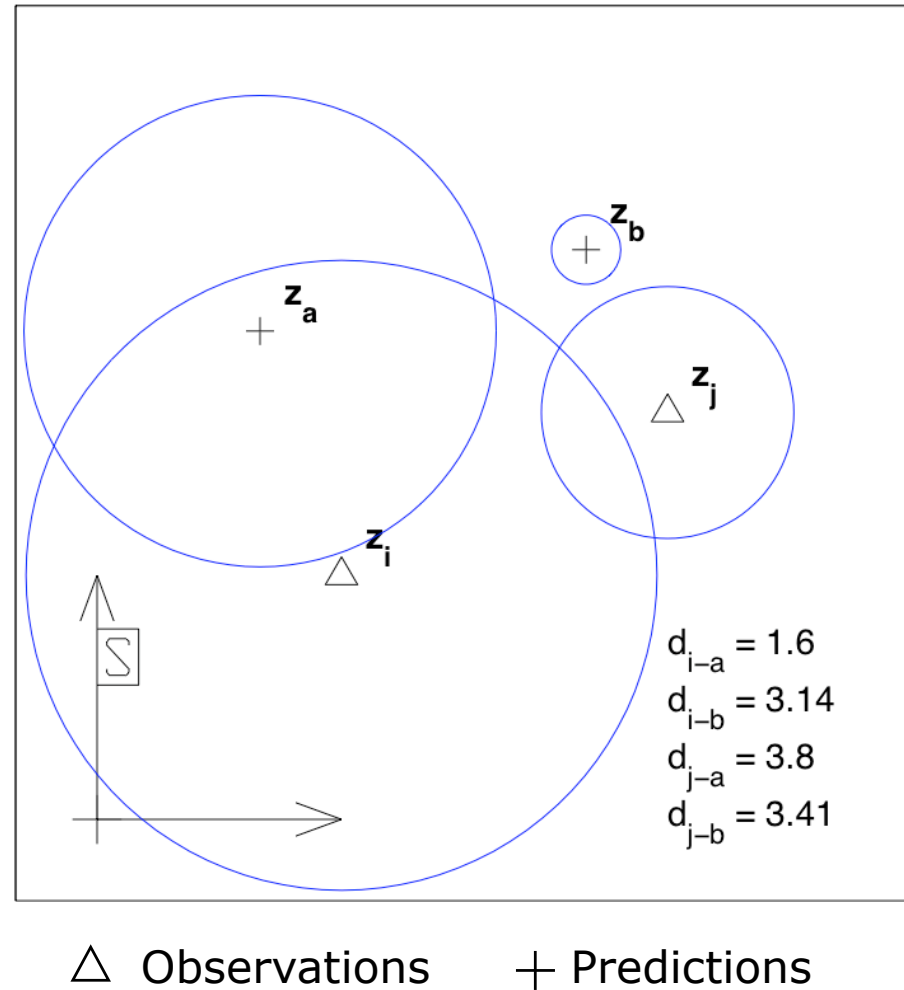
Validation Gate

Mahalanobis distance
with **diagonal** covariance
matrices

Takes into account:

- ✓ Position
- ✓ Uncertainty
- ✗ Correlations

→ Now, i-b is “closer”
than j-b



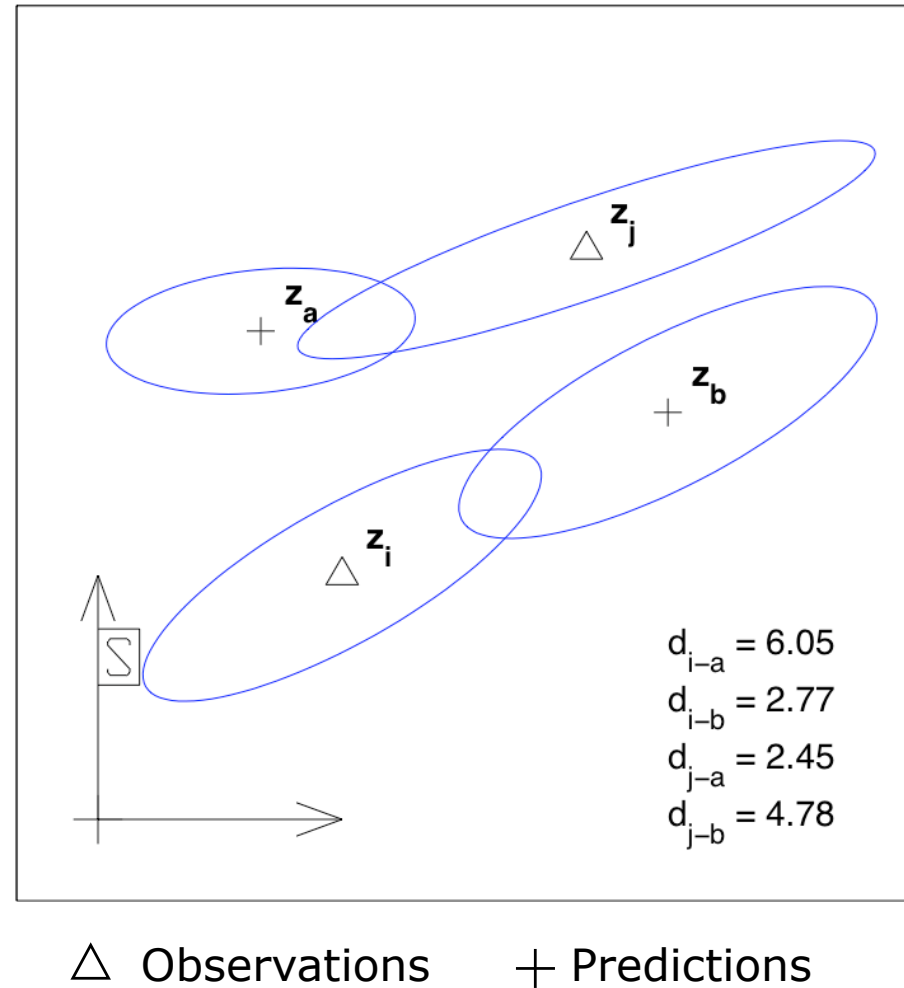
Validation Gate

Mahalanobis distance

Takes into account:

- ✓ Position
- ✓ Uncertainty
- ✓ Correlations

→ It's actually i-b and j-a that belong together!



False Alarms

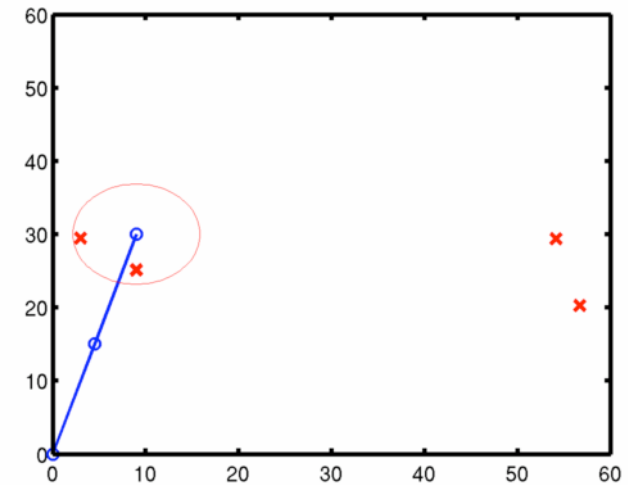
- False alarms are **false positives**
- They can come from sensor imperfections or detector failures
- They raise the two questions:

What is actually inside my **validation gate**?

- The real measurement or
- a false alarm?

How to **model false alarms**?

- Uniform over sensor space
- Independent across time

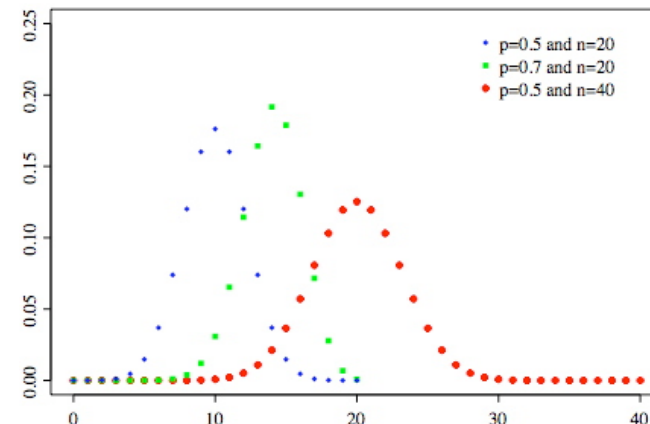


False Alarm Model

- Assume (temporarily) that the sensor field of view V is discretized into N discrete cells, c_i , $i = 1, \dots, N$
- In each cell, false alarms occur with probability P_F
- Assume independence across cells
- The occurrence of false alarms is a Bernoulli process (flipping an unfair coin) with probability $p = P_F$
- Then, the number of false alarms m_F follows a **Binomial distribution**

$$P(K = m_F) = \binom{N}{m_F} p^{m_F} (1 - p)^{N - m_F}$$

with expected value Np



False Alarm Model

- Let the spatial density λ be the number of false alarms over space

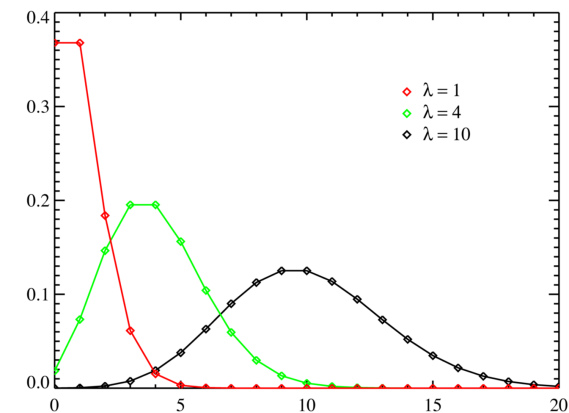
$$\lambda = \frac{Np}{V} \quad [\text{occurrences per m}^2]$$

- Let now $N \rightarrow \infty$, that is, we reduce the cell size until the continuous case. Then the Binomial becomes a Poisson distribution with

$$\mu_F(m_F) = e^{-\lambda V} \frac{(\lambda V)^{m_F}}{m_F!}$$

- The **measurement likelihood** of false alarms is assumed to be uniform,

$$p(z|z \text{ is a false alarm}) = \frac{1}{V}$$



Single Target Data Association

Let us consider

- A **single** target to track
- Track already initialized
- Detection probability < 1
- False alarm probability > 0

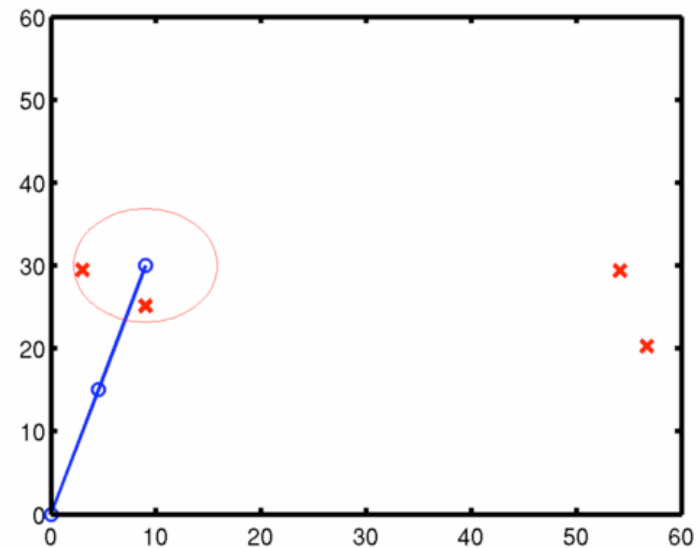
Data association approaches

Non-Bayesian:

- Nearest neighbor (NN)
- Track splitting filter

Bayesian:

- Probabilistic Data Association Filter (PDAF)



Single Target DA: NN

Nearest Neighbor filter (NN)

1. Compute Mahalanobis distance to all measurements
 2. Accept the **closest measurement**
 3. Update the track as if it were the correct measurement
- Problem: with some probability the selected measurement is not the correct one. This can lead to filter divergence (covariances collapse regardless)
 - **Conservative NN variant:**
Do not associate in case of ambiguities

Single Target DA: PDAF

Probabilistic Data Association filter (PDAF)

- Integrates **all** measurements in the validation gate

- Conditioning the update on

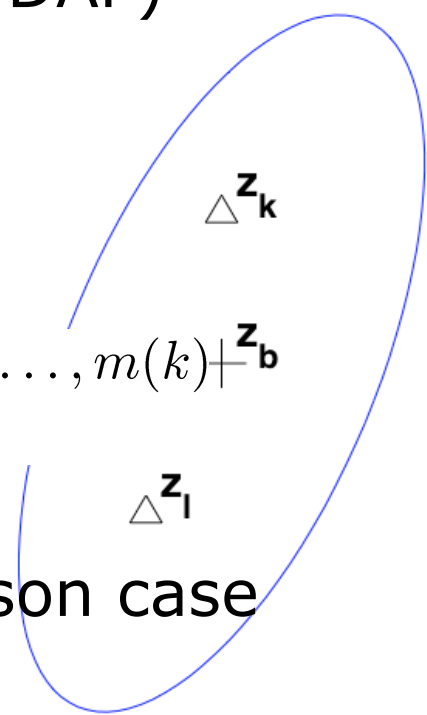
$$\theta_i(k) = \begin{cases} z_i(k) \text{ is the correct measurement} & i = 1, \dots, m(k) \\ \text{no correct measurement is present} & i = 0 \end{cases} \quad \begin{matrix} \mathbf{z}_k \\ \mathbf{z}_b \end{matrix}$$

- With probability $\beta_i \triangleq P(\theta_i | Z^k)$ for the Poisson case

$$\beta_i(k) = \begin{cases} \frac{e_i}{b + \sum_{j=1}^{\mu_F} e_j} & i = 1, \dots, m(k) \\ \frac{b}{b + \sum_{j=1}^{\mu_F} e_j} & i = 0 \end{cases}$$

$$e_i = \mu_F(m(k) - 1) \cdot P_D P_G \cdot P_G^{-1} \mathcal{N}(\nu_i(k); 0, \hat{S}(k))$$

$$b = \mu_F(m(k))(1 - P_D P_G)$$



Single Target DA: PDAF

- Uses all the measurements in the validation area
 - Conditioning the update on

$$\theta_i(k) = \begin{cases} z_i(k) \text{ is the correct measurement} & i = 1, \dots, m(k) \\ \text{no correct measurement is present} & i = 0 \end{cases}$$

- With probability $\beta_i \triangleq P(\theta_i | Z^k)$ for the Poisson case

$$\beta_i(k) = \begin{cases} \frac{e_i}{b + \sum_{j=1}^m \mu_F} & i = 1, \dots, m(k) \\ \frac{b}{b + \sum_{j=1}^m \mu_F} & i = 0 \end{cases}$$

$$e_i = \mu_F(m(k) - 1) \cdot P_D P_G \cdot P_G^{-1} \mathcal{N}(\nu_i(k); 0, \hat{S}(k))$$

$$b = \mu_F(m(k))(1 - P_D P_G)$$

Single Target DA: PDAF

- State update

$$\hat{x}(k|k) = \hat{x}(k|k-1) + K(k)\nu(k)$$

- With the combined innovation

$$\nu(k) = \sum_{i=1}^N \beta_i(k) \nu_i(k)$$

- Covariance update

$$P(k|k) = \beta_0(k)P(k|k-1) + (1 - \beta_0(k))P(k|k) + \tilde{P}(k)$$

- With the spread of innovations

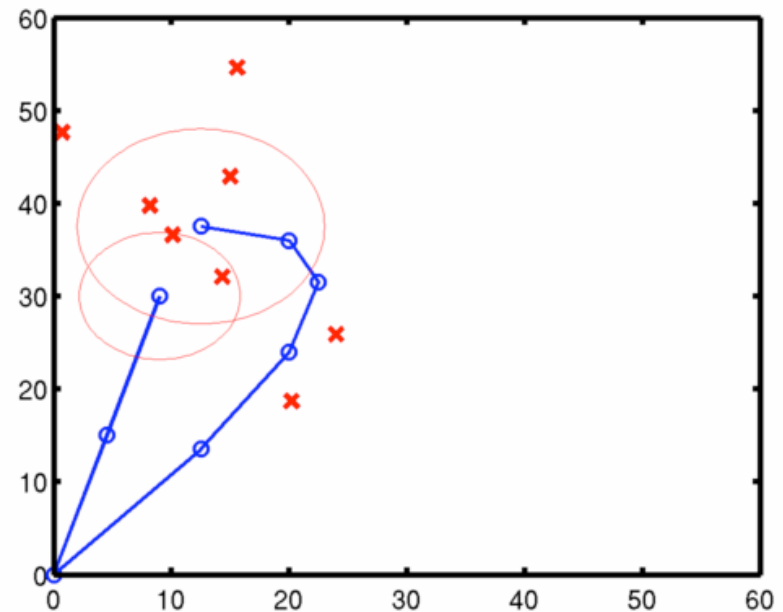
$$\tilde{P}(k) = K(k) \left[\sum_{i=1}^N \beta_i(k) \nu_i(k) \nu_i(k)^T - \nu(k) \nu(k)^T \right] K(k)^T$$

Single Target DA: Summary

- **Nearest Neighbor filter (NN)**
 - Simple to implement
 - Can integrate wrong measurements (false alarms), and thus, produce overconfident estimates
 - Good if prediction and measurement models are accurate
- **Probabilistic Data Association filter (PDAF)**
 - A bit more involved to implement
 - Provide conservative estimates
 - Good in presence of high clutter and noisy models

Multi-Target Data Association

- Multiple targets to track
 - Tracks already initialized
 - Detection probability < 1
 - False alarm probability > 0
- Non Bayesian approaches
 - Nearest neighbor
 - Interpretation tree
 - Joint compatibility (JCBB)
- Bayesian approaches
 - JPDAF
 - MHT
 - MCMC



Multi-Target DA: NN

- Build the assignment matrix $A = [d_{ij}^2]$

$$d_{ij}^2 = \nu_{ij}(k)^T S_j^{-1}(k) \nu_{ij}(k)$$

- Iterate
 - Find the minimum cost assignment in A
 - Remove the row and column of that assignment
- Check if assignment are in the validation regions
 - Unassociated tracks can be used for track deletion
 - Unassociated measurements can be used for track creation
- Problem: It's not a global minimum
- Conservative NN variant: no association in case of ambiguities

Multi-Target DA: Global NN

- Build the assignment matrix $A = [d_{ij}^2]$

$$d_{ij}^2 = \nu_{ij}(k)^T S_j^{-1}(k) \nu_{ij}(k)$$

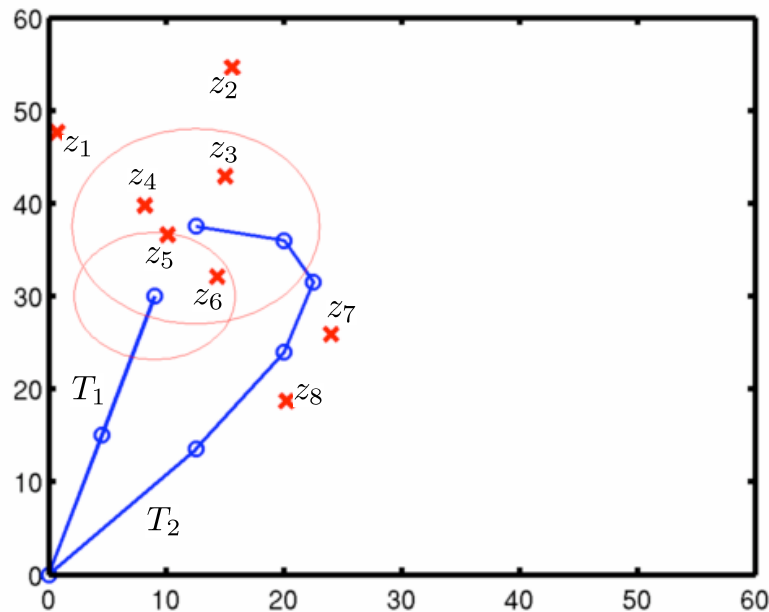
- Solve the linear assignment problem

$$\min \sum d_{ij}^2 \cdot x_{ij} \quad x_{ij} \in \{0, 1\}$$

$$\sum_i x_{ij} = 1 \quad \sum_j x_{ij} = 1$$

- **Hungarian** method (blow up to square matrix)
 - **Munkres** algorithm for rectangular matrices
 - Finds **global** cost minimum!
- Check if assignments are in the validation gate
- Performs DA jointly!

Assignment Matrix Example



- Rectangular

$$A = \begin{bmatrix} d_{11}^2 & d_{12}^2 & d_{13}^2 & d_{14}^2 & d_{15}^2 & d_{16}^2 & d_{17}^2 & d_{18}^2 \\ d_{21}^2 & d_{22}^2 & d_{23}^2 & d_{24}^2 & d_{25}^2 & d_{26}^2 & d_{27}^2 & d_{28}^2 \end{bmatrix}$$

- Square

$$A = \begin{bmatrix} d_{11}^2 & d_{12}^2 & d_{13}^2 & d_{14}^2 & d_{15}^2 & d_{16}^2 & d_{17}^2 & d_{18}^2 \\ d_{21}^2 & d_{22}^2 & d_{23}^2 & d_{24}^2 & d_{25}^2 & d_{26}^2 & d_{27}^2 & d_{28}^2 \\ p_F & p_F & p_F & p_F & p_F & p_F & p_F & p_F \\ p_F & p_F & p_F & p_F & p_F & p_F & p_F & p_F \\ p_F & p_F & p_F & p_F & p_F & p_F & p_F & p_F \\ p_F & p_F & p_F & p_F & p_F & p_F & p_F & p_F \\ p_F & p_F & p_F & p_F & p_F & p_F & p_F & p_F \\ p_F & p_F & p_F & p_F & p_F & p_F & p_F & p_F \end{bmatrix}$$

- Entries

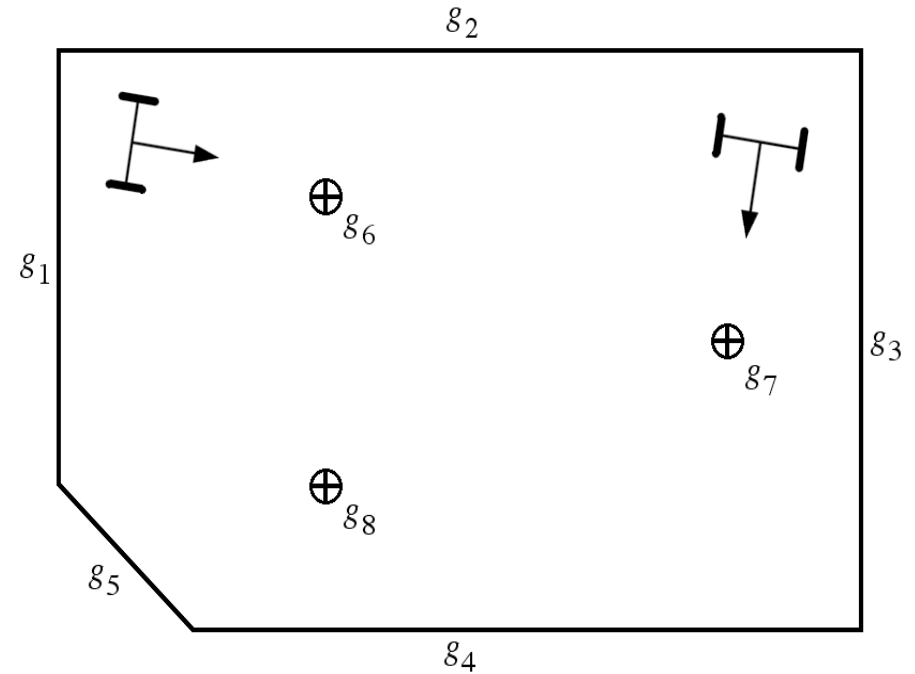
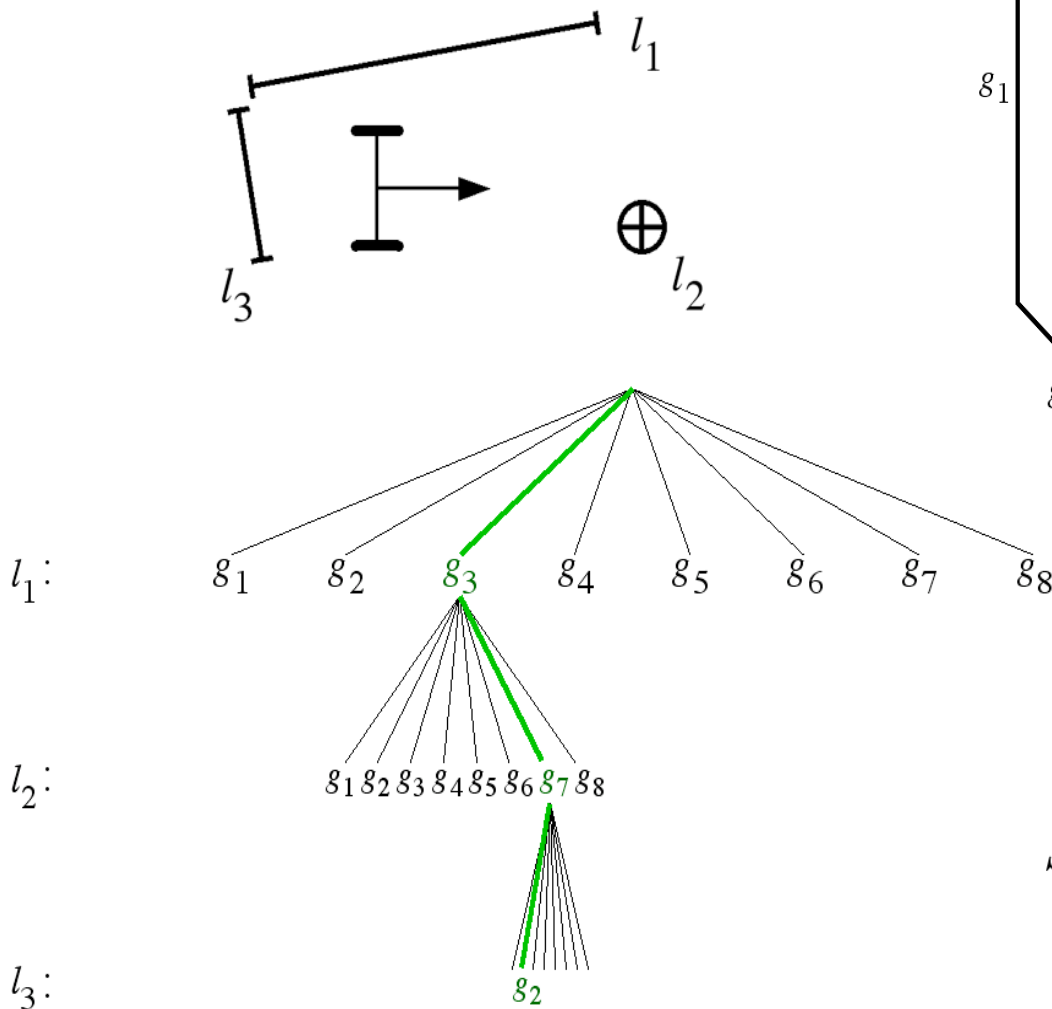
- $d_{ij}^2 = \nu_{ij}(k)^T S_j^{-1}(k) \nu_{ij}(k)$
- False alarm probability

GNN vs. Interpretation Tree

- A solution to GNN in presence of constraints
 - Introduced in [Grimson 87], used in feature-based SLAM
- Main idea: consider all possible association among measurements and tracks
 - The association are built over a tree representation
 - A Depth-first visit on the tree is performed
 - Constraints are used to prune the tree
- Worst case: exponential complexity
 - The whole number of associations is $(\#T + 1)^{\#m}$

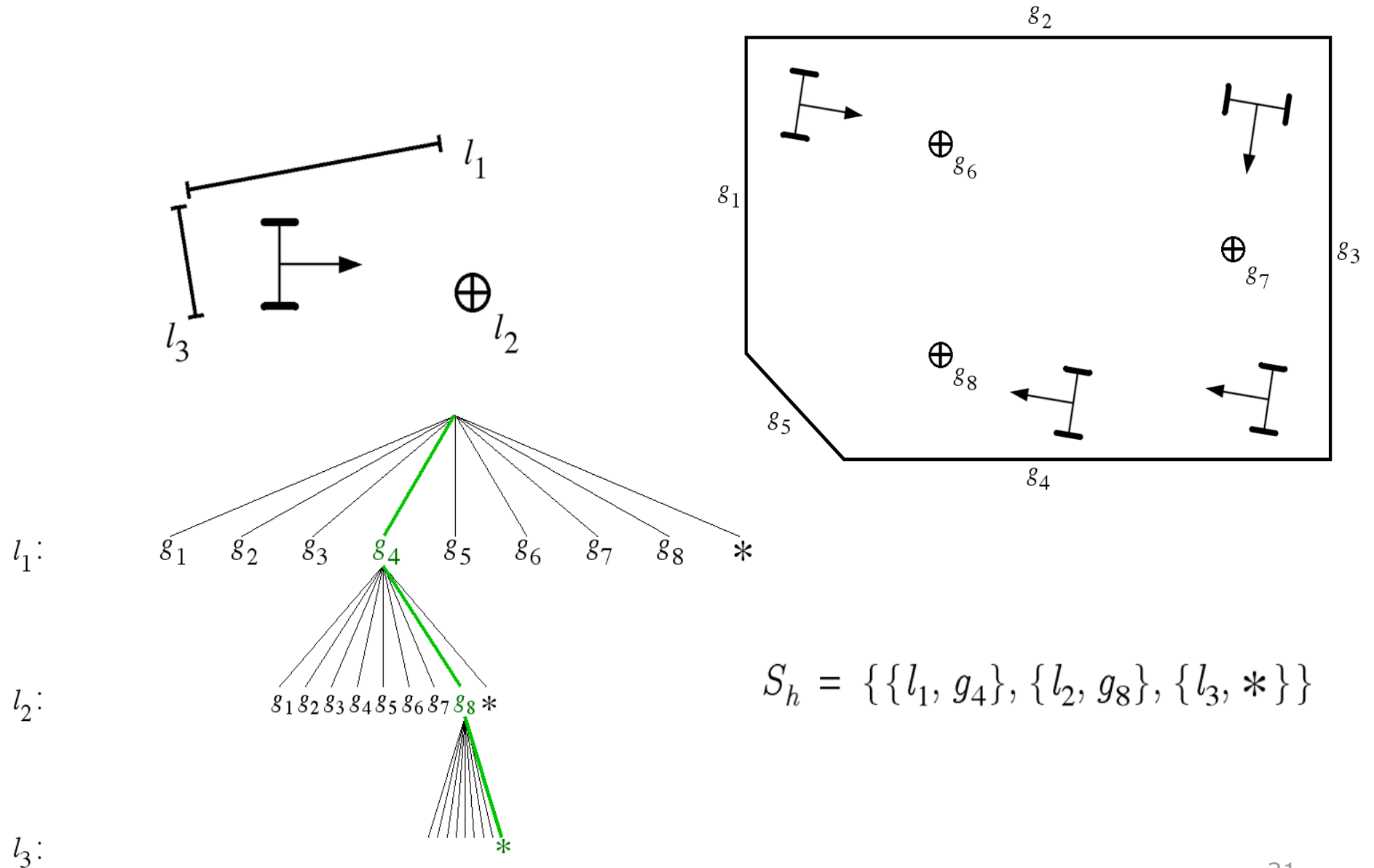
Interpretation Tree: SLAM

Interpretation tree



$$S_{h_2} = \{\{l_1, g_3\}, \{l_2, g_7\}, \{l_3, g_2\}\}$$

Interpretation Tree: SLAM



GNN Comparison

Assignment solver

- Pros
 - Fast: polynomial
 - Libraries available
 - Extension to k-best
- Cons
 - Only linear constraints
 - Blow-up of cost matrix

Interpretation tree

- Pros
 - General constraints
 - Extensive search
 - Provide k-best
- Cons
 - Slow: exponential

Interpretation trees are good when several constraints are available, in other cases assignment solver are preferable

Joint Compatibility

- Individual compatibility (e.g. independent tracks)
 - One measurement integration influences only one track
 - One measurement per track assumption
 - Typical of target tracking
- Joint compatibility (e.g. correlated tracks)
 - One measurement integration influences several track
 - Multiple measurement per track association
 - Typical of localization and SLAM

Joint Compatibility

- Given the joint hypothesis $\mathcal{H} = \{j_1, \dots, j_m\}$
 - the k-th measurement is associated with track $j_k = t$
- And the joint measurement function

$$\begin{aligned}\mathbf{z}_{\mathcal{H}} &= \mathbf{h}_{\mathcal{H}}(\mathbf{x}) + \mathbf{w} \\ \mathbf{h}_{\mathcal{H}} &= \begin{bmatrix} \mathbf{h}_{ij_1} \\ \vdots \\ \mathbf{h}_{ij_m} \end{bmatrix}\end{aligned}$$

- The hypothesis is jointly compatible if

$$\begin{aligned}D_{\mathcal{H}}^2 &= (\mathbf{z}_{\mathcal{H}} - \mathbf{h}_{\mathcal{H}}(\hat{\mathbf{x}}))^T C_{\mathcal{H}}^{-1} (\mathbf{z}_{\mathcal{H}} - \mathbf{h}_{\mathcal{H}}(\hat{\mathbf{x}})) \leq \chi_{d,\alpha}^2 \\ C_{\mathcal{H}} &= H_{\mathcal{H}} \hat{P} H_{\mathcal{H}}^T + R_{\mathcal{H}}\end{aligned}$$

Joint Compatibility – JCBB

- Joint compatibility branch and bound
- Initialize with empty hypothesis and first obs.
- For all tracks
 - If is individually and jointly compatible assign and recursively call JCBB
 - Otherwise consider it a false alarm

```
procedure JCBB (H, i):  
  -- H : current hypothesis  
  -- i : observation to be matched  
  if i > m  
    if pairings(H) > pairings(Best)  
      Best = H  
    fi  
  else  
    for j in {1...n}  
      if unary(i, j)  
        ∧ joint_compatibility([H j])  
        JCBB([H j], i + 1)  
      fi  
    rof  
    if pairings(H) + m - i > pairings(Best)  
      JCBB([H 0], i + 1)  
    fi  
  fi
```

[Neira et al.'03]

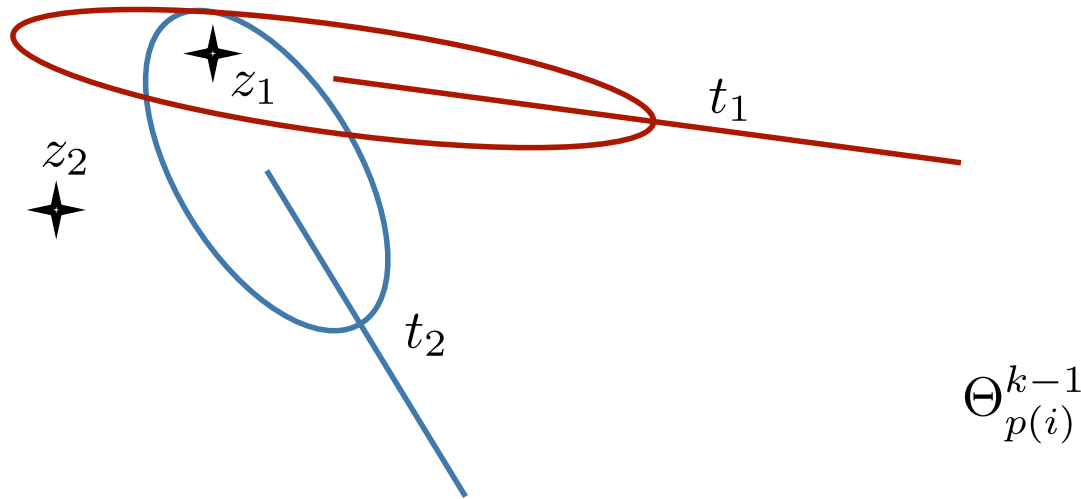
Multi-Target DA: MHT

- Reason about the associations of sequences measurements with tracks and false alarm
- Evaluate the probability of association hypotheses
- Optimal Bayesian solution
- Algorithm
 - State and measurement prediction
 - Hypotheses generation
 - Hypotheses probability evaluation
 - State update
 - Hypotheses management (i.e. pruning, elimination, creation)
- Exponential complexity of the full solution
 - Pruning strategies
 - K-best hypotheses

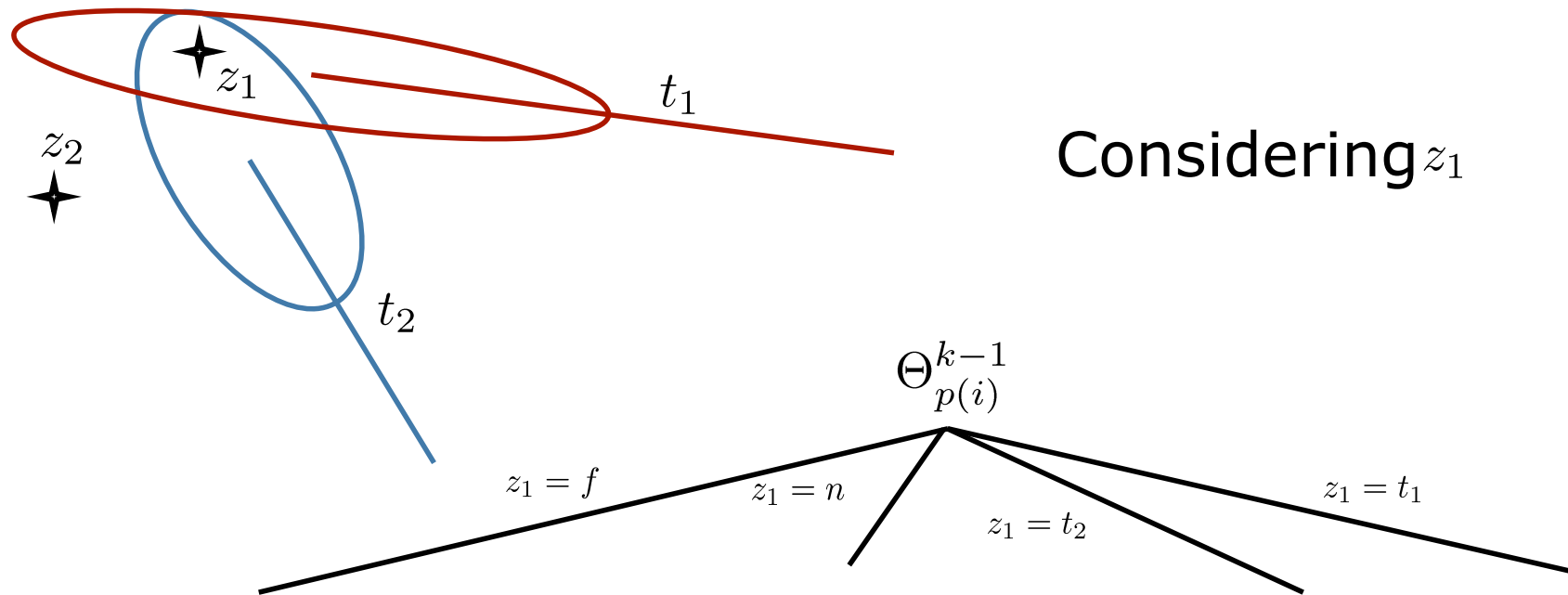
MHT: Hypothesis Generation

- A hypothesis $\Theta_i^k = \{\Theta_{p(i)}^{k-1}, \theta_{c(i)}(k)\}$ at time k is a history of assignment sets $\theta = \{\theta_{obs}, \theta_{track}\}$ to time k
 - $\theta_{obs} = \{z_1, \dots, z_{m(k)}\}$ is a set of measurement associations, where a measurement is either associated to track $z_i = t$, treated as a new track $z_i = n$ or as a false alarm $z_i = f$
 - $\theta_{track} = \{l_1, \dots, l_{t(k)}\}$ is a set of track label, where a track can be matched $l_i = m$, occluded $l_i = o$ or deleted $l_i = d$
- Hypotheses are generated recursively in a tree-based structure
 - Unlikely branches are avoided by validation gating
 - Exponential growth of the trees
 - Only a subset of hypotheses are generated in practice

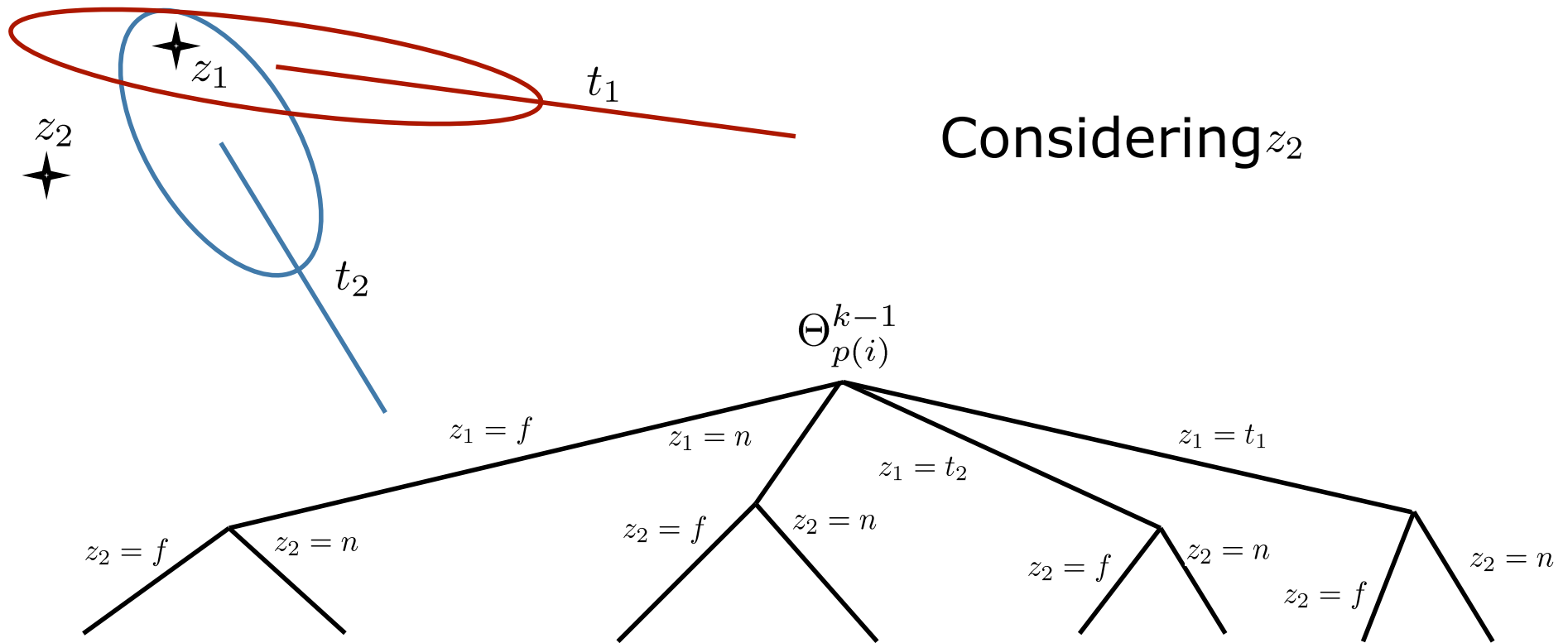
MHT: Hypothesis Generation



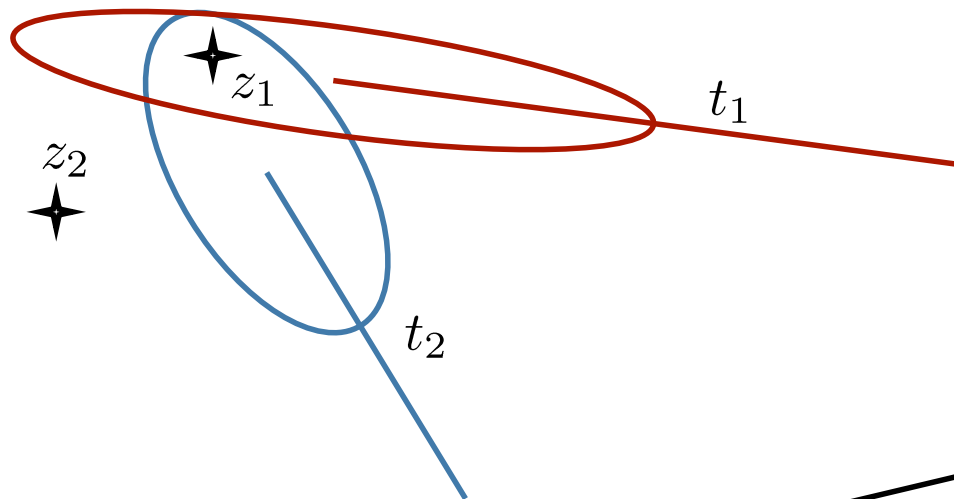
MHT: Hypothesis Generation



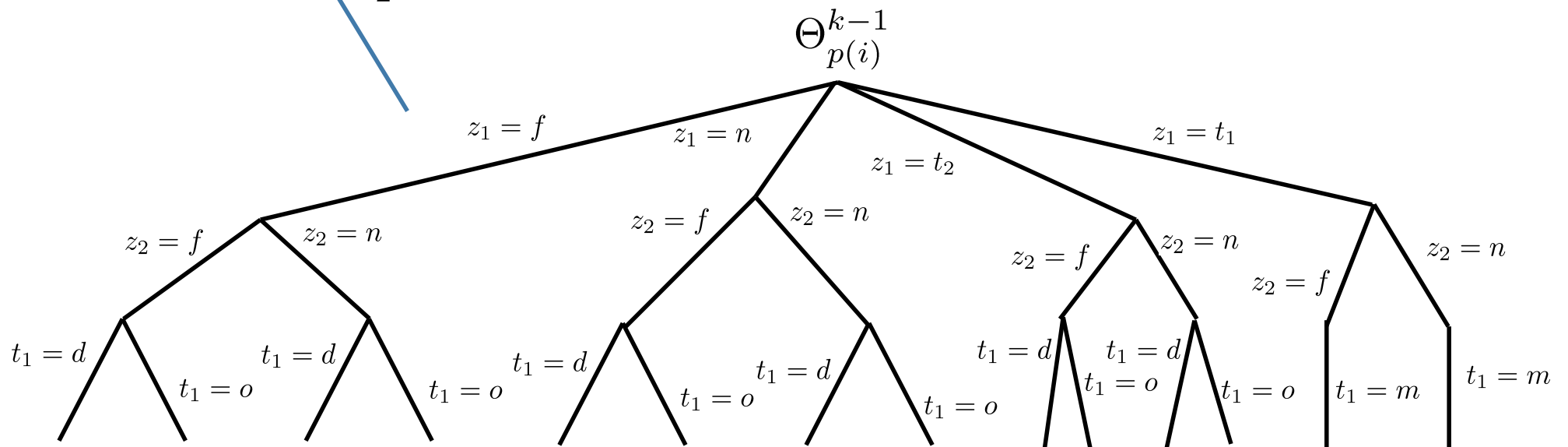
MHT: Hypothesis Generation



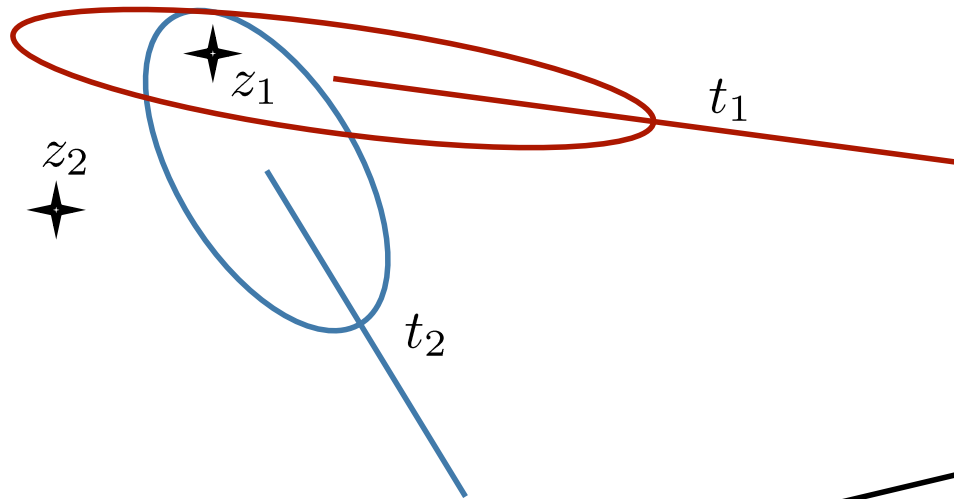
MHT: Hypothesis Generation



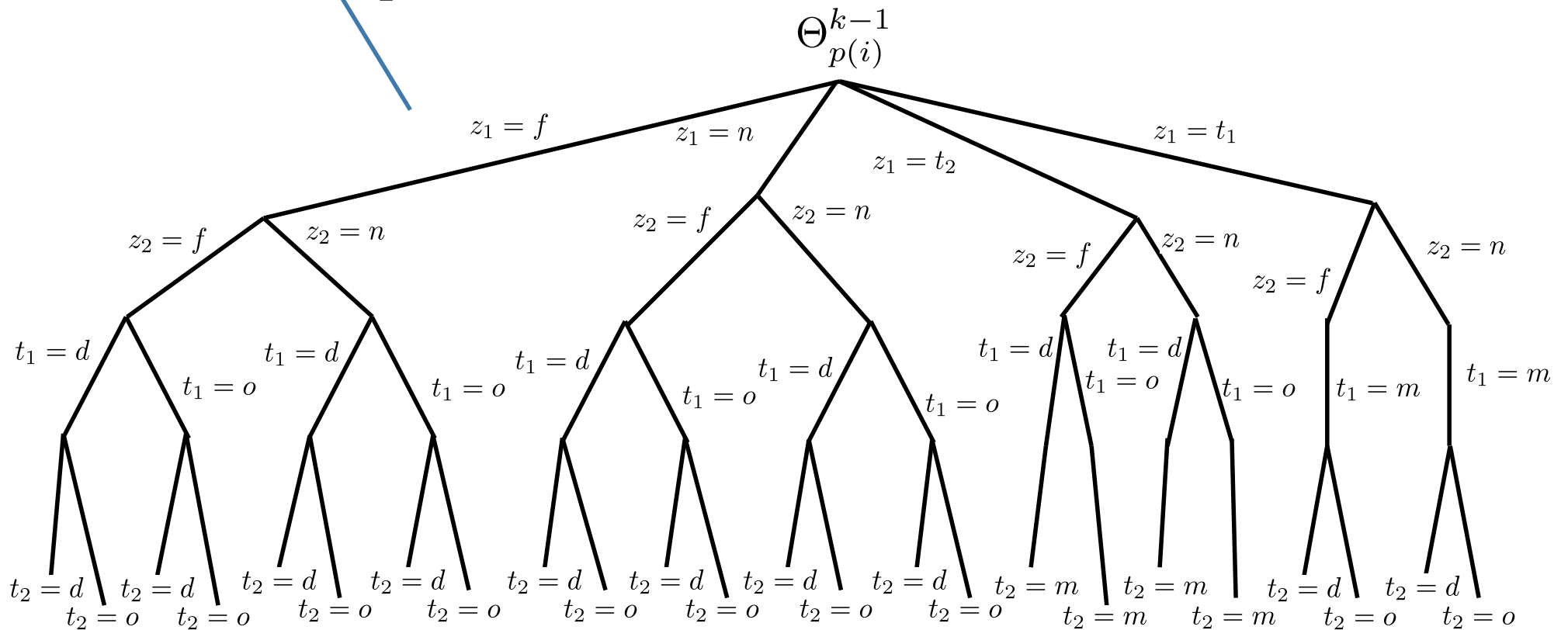
Considering t_1



MHT: Hypothesis Generation



Considering t_2



MHT: Hypothesis Evaluation

- The probability of an hypothesis $\Theta_i^k = \{\Theta_{p(i)}^{k-1}, \theta_{c(i)}(k)\}$ can be calculated using Bayes rules

$$P(\Theta_i^k | Z^k) = P(\Theta_{p(i)}^{k-1}, \theta_{c(i)}(k) | Z^k) =$$
$$\frac{1}{\eta} \cdot \underbrace{p(Z(k) | \Theta_{p(i)}^{k-1}, \theta_{c(i)}(k), Z^{k-1})}_{\text{Likelihood}} \cdot \underbrace{P(\theta_{c(i)}(k) | \Theta_{p(i)}^{k-1}, Z^k)}_{\text{Assignment probability}} \cdot \underbrace{P(\Theta_{p(i)}^{k-1} | Z^{k-1})}_{\text{Prior}}$$

MHT: Hypothesis Evaluation

- Likelihood

$$p(Z(k)|\Theta^{k-1}, \theta(k), Z^{k-1}) = \prod_{l=1}^{m(k)} p(z_l(k)|\Theta^{k-1}, \theta(k), Z^{k-1})$$

- Case 1: associated with track t

$$p(z_l(k)|\Theta^{k-1}, \theta(k), Z^{k-1}) = \mathcal{N}(z_l(k); \hat{z}_t(k|k-1), S_t(k))$$

- Case 2: false alarm

$$p(z_l(k)|\Theta^{k-1}, \theta(k), Z^{k-1}) = V^{-1}$$

- Case 3: new track

$$p(z_l(k)|\Theta^{k-1}, \theta(k), Z^{k-1}) = V^{-1}$$

MHT: Hypothesis Evaluation

- Assignment probability

$$P(\theta(k)|\Theta^{k-1}, Z^k) = P(\theta(k)|N_M, N_O, N_D, N_N, N_F) \cdot P(N_M, N_O, N_D, N_N, N_F|\theta(k), \Theta^{k-1})$$

- $P(N_M, N_O, N_D, N_N, N_F|\theta(k), \Theta^{k-1})$ is the probability of having N_M matched tracks, N_O occluded tracks, N_D deleted tracks, N_F false alarm and N_N new tracks.
- $P(\theta(k)|N_M, N_O, N_D, N_N, N_F)$ is the probability of a possible configuration $\theta(k)$ given the number of events defined before

MHT: Hypothesis Evaluation

- Assignment probability 1: $P(N_M, N_O, N_D, N_N, N_F | \theta(k), \Theta^{k-1})$
 - Assuming a multinomial distribution for track labels

$$P(N_M, N_O, N_D | \theta(k), \Theta^{k-1}) = \frac{N_T!}{N_M! N_O! N_D!} p_M^{N_M} p_O^{N_O} p_D^{N_D}$$

- Assuming a Poisson distribution for new tracks

$$P(N_N | \theta(k), \Theta^{k-1}) = \frac{(V\lambda_N)^{N_N} e^{-V\lambda_N}}{N_N!}$$

- Assuming a Poisson distribution for false alarm

$$P(N_F | \theta(k), \Theta^{k-1}) = \frac{(V\lambda_F)^{N_F} e^{-V\lambda_F}}{N_F!}$$

- We obtain

$$P(\cdot) = \frac{N_T! (e^{-V\lambda_N}) (e^{-V\lambda_F})}{N_N! N_F! N_M! N_O! N_D!} (V\lambda_N)^{N_N} (V\lambda_F)^{N_F} p_M^{N_M} p_O^{N_O} p_D^{N_D}$$

MHT: Hypothesis Evaluation

- Assignment probability 2: $P(\theta(k)|N_M, N_O, N_D, N_N, N_F)$

- The possible choices of $m(k)$ taken as matched tracks

$$\binom{m(k)}{N_M} Perm(N_M, N_T) = \frac{m(k)!}{N_M!(m(k) - N_M)!} \frac{N_T!}{(N_T - N_M)!}$$

- The combinations of $m(k) - N_M$ taken as new tracks or false alarms

$$\binom{m(k) - N_M}{N_N} \binom{m(k) - N_M - N_N}{N_F = 1} = \frac{(m(k) - N_M)!}{N_N!(m(k) - N_M - N_N)!}$$

- The combinations of $N_T - N_M$ taken as occluded or deleted

$$\binom{N_T - N_M}{N_O} \binom{N_T - N_M - N_O}{N_D = 1} = \frac{(N_T - N_M)!}{N_O!(N_T - N_M - N_O)!}$$

- The probability is 1 over all the possible choices

$$\left[\frac{m(k)!}{N_M!(m(k) - N_M)!} \frac{N_T!}{(N_T - N_M)!} \frac{(m(k) - N_M)!}{N_N!(m(k) - N_M - N_N)!} \frac{(N_T - N_M)!}{N_O!(N_T - N_M - N_O)!} \right]^{-1}$$

MHT: Hypothesis Evaluation

- Assignment probability 2: $P(\theta(k)|N_M, N_O, N_D, N_N, N_F)$

- The possible choices of $m(k)$ taken as matched tracks

$$\binom{m(k)}{N_M} Perm(N_M, N_T) = \frac{m(k)!}{N_M!(m(k) - N_M)!} \frac{N_T!}{(N_T - N_M)!}$$

- The combinations of $m(k) - N_M$ taken as new tracks or false alarms

$$\binom{m(k) - N_M}{N_N} \binom{m(k) - N_M - N_N}{N_F = 1} = \frac{(m(k) - N_M)!}{N_N!(m(k) - N_M - N_N)!}$$

- The combinations of $N_T - N_M$ taken as occluded or deleted

$$\binom{N_T - N_M}{N_O} \binom{N_T - N_M - N_O}{N_D = 1} = \frac{(N_T - N_M)!}{N_O!(N_T - N_M - N_O)!}$$

- The probability is 1 over all the possible choices

$$\left[\frac{m(k)!}{N_M!(m(k) - N_M)!} \frac{N_T!}{(N_T - N_M)!} \frac{(m(k) - N_M)!}{N_N! \underbrace{(m(k) - N_M - N_N)}_{N_D}} \frac{(N_T - N_M)!}{N_O! \underbrace{(N_T - N_M - N_O)}_{N_F}} \right]^{-1}$$

MHT: Hypothesis Evaluation

- Assignment probability 2: $P(\theta(k)|N_M, N_O, N_D, N_N, N_F)$

- The possible choices of $m(k)$ taken as matched tracks

$$\binom{m(k)}{N_M} Perm(N_M, N_T) = \frac{m(k)!}{N_M!(m(k) - N_M)!} \frac{N_T!}{(N_T - N_M)!}$$

- The combinations of $m(k) - N_M$ taken as new tracks or false alarms

$$\binom{m(k) - N_M}{N_N} \binom{m(k) - N_M - N_N}{N_F = 1} = \frac{(m(k) - N_M)!}{N_N!(m(k) - N_M - N_N)!}$$

- The combinations of $N_T - N_M$ taken as occluded or deleted

$$\binom{N_T - N_M}{N_O} \binom{N_T - N_M - N_O}{N_D = 1} = \frac{(N_T - N_M)!}{N_O!(N_T - N_M - N_O)!}$$

- The probability is 1 over all the possible choices

$$P(\theta(k)|N_M, N_O, N_D, N_N, N_F) = \frac{N_M!N_N!N_F!N_O!N_D!}{m(k)!N_T!}$$

MHT: Hypothesis Evaluation

- Assignment probability

$$P(\theta(k)|\Theta^{k-1}, Z^k) = P(\theta(k)|N_M, N_O, N_D, N_N, N_F) \cdot \\ \cdot P(N_M, N_O, N_D, N_N, N_F|\theta(k), \Theta^{k-1})$$

- Combining everything together we have

$$P(\theta(k)|\Theta^{k-1}, Z^k) = \\ = \frac{N_T!(e^{-V\lambda_N})(e^{-V\lambda_F})}{N_N!N_F!N_M!N_O!N_D!} (V\lambda_N)^{N_N} (V\lambda_F)^{N_F} p_M^{N_M} p_O^{N_O} p_D^{N_D} \frac{N_M!N_N!N_F!N_O!N_D!}{m(k)!N_T!}$$

MHT: Hypothesis Evaluation

- Assignment probability

$$P(\theta(k)|\Theta^{k-1}, Z^k) = P(\theta(k)|N_M, N_O, N_D, N_N, N_F) \cdot P(N_M, N_O, N_D, N_N, N_F|\theta(k), \Theta^{k-1})$$

- Combining everything together we have

$$P(\theta(k)|\Theta^{k-1}, Z^k) = \frac{\cancel{N_F!} (e^{-V\lambda_N})(e^{-V\lambda_F})}{\cancel{N_N!N_F!N_M!N_O!N_D!}} (V\lambda_N)^{N_N} (V\lambda_F)^{N_F} p_M^{N_M} p_O^{N_O} p_D^{N_D} \frac{\cancel{N_M!N_N!N_F!N_O!N_D!}}{m(k)! \cancel{N_T!}}$$

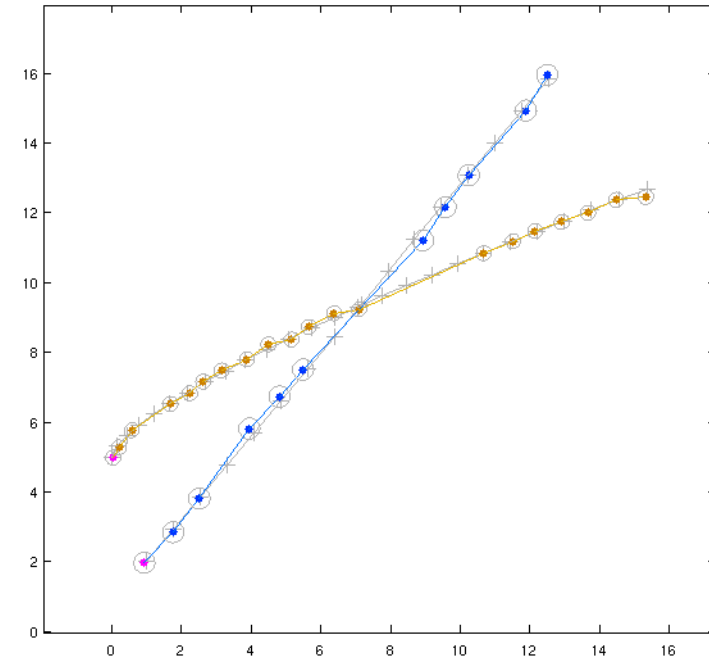
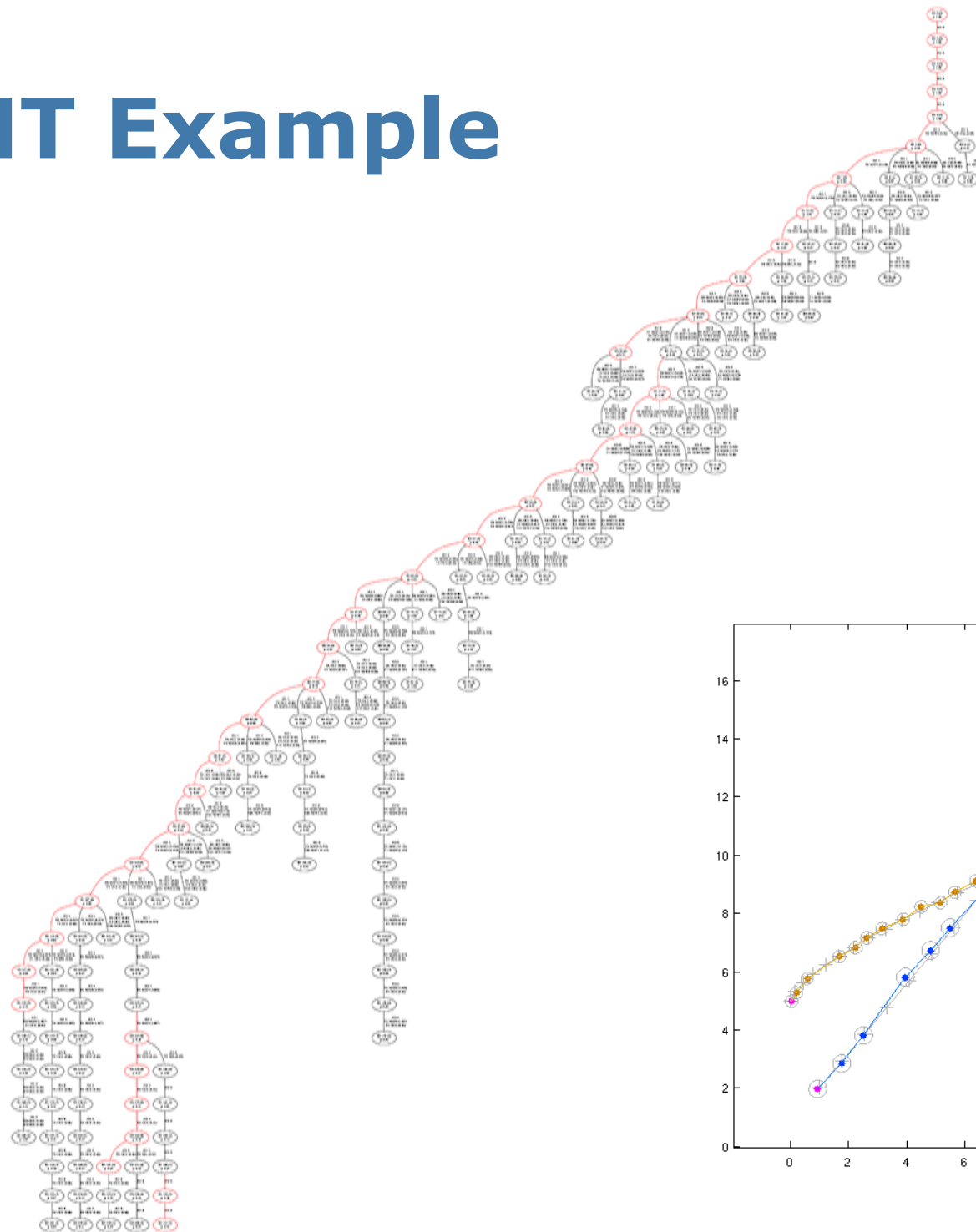
- Simplifying the expression we obtain

$$P(\theta(k)|\Theta^{k-1}, Z^k) = \frac{(e^{-V\lambda_N})(e^{-V\lambda_F})}{m(k)!} (V\lambda_N)^{N_N} (V\lambda_F)^{N_F} p_M^{N_M} p_O^{N_O} p_D^{N_D}$$

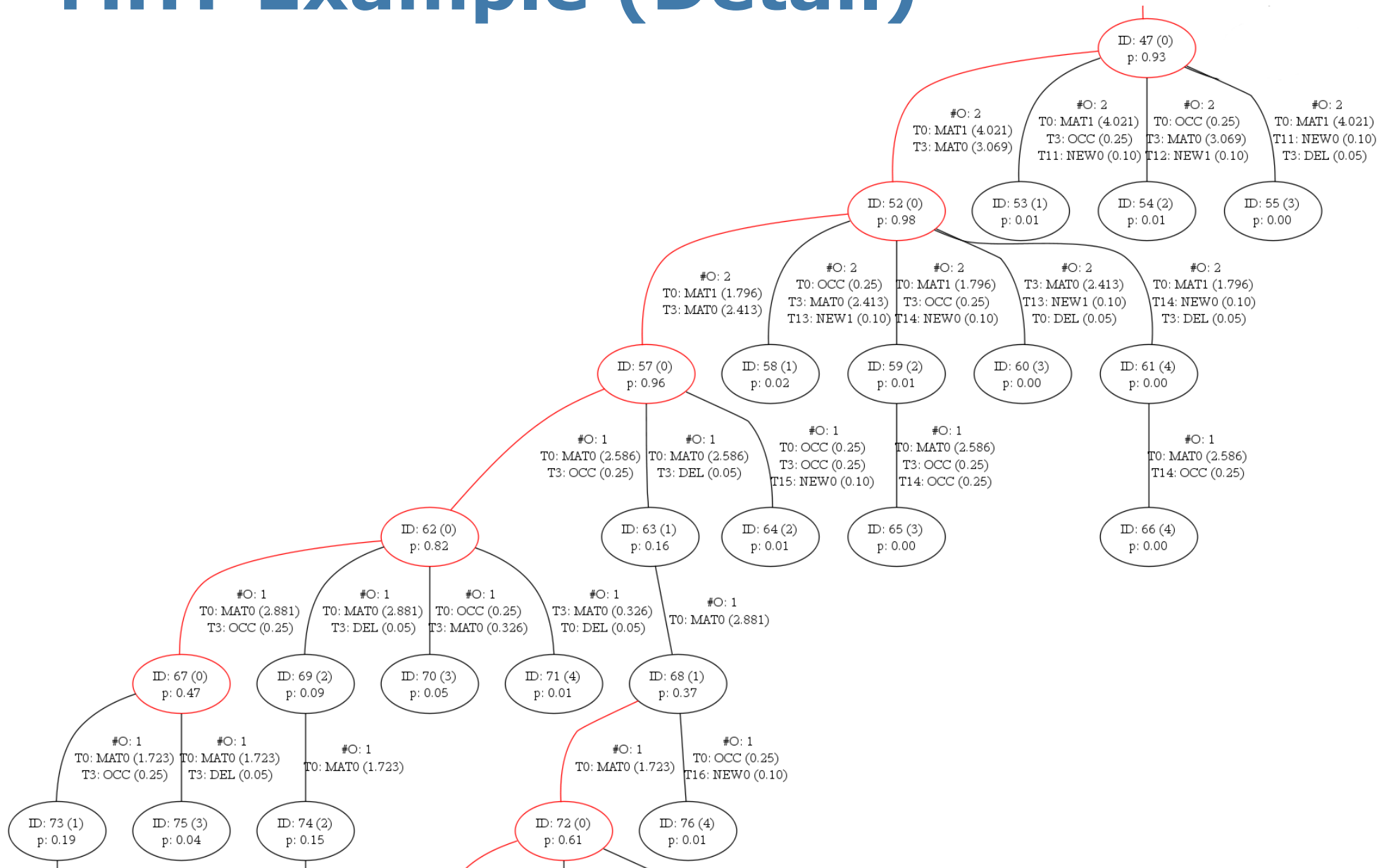
MHT Approximations

- Spatially disjoint hypothesis trees
 - Tracks can be partitioned in clusters
 - A separate tree is grown for each cluster
- K-Best hypothesis tree
 - Directly generate the k-best hypothesis
 - Generation and evaluation are integrated in a single step
 - Use Murty algorithm and a linear assignment solver
- N-Scan back pruning
 - Ambiguities are supposed to be resolved after N steps
 - Children at step $k+N$ give the prob. of parents at step k
 - Keep only the most probable branch

MHT Example



MHT Example (Detail)



Multi-Target DA: Summary

- **Nearest Neighbor filters** (NN and GNN)
 - Simple to implement
 - NN: Good if tracks are well separated and not noisy
 - NN+GNN: No integration over time
- **Interpretation tree**
 - More involved to implement
 - Good in case of general constraints among associations
- **MHT**
 - Fully Bayesian
 - Most general framework for multiple targets
 - Complex and expensive
 - Only approximations are practically implemented