

Teaser application 2: Place recognition in changing environments

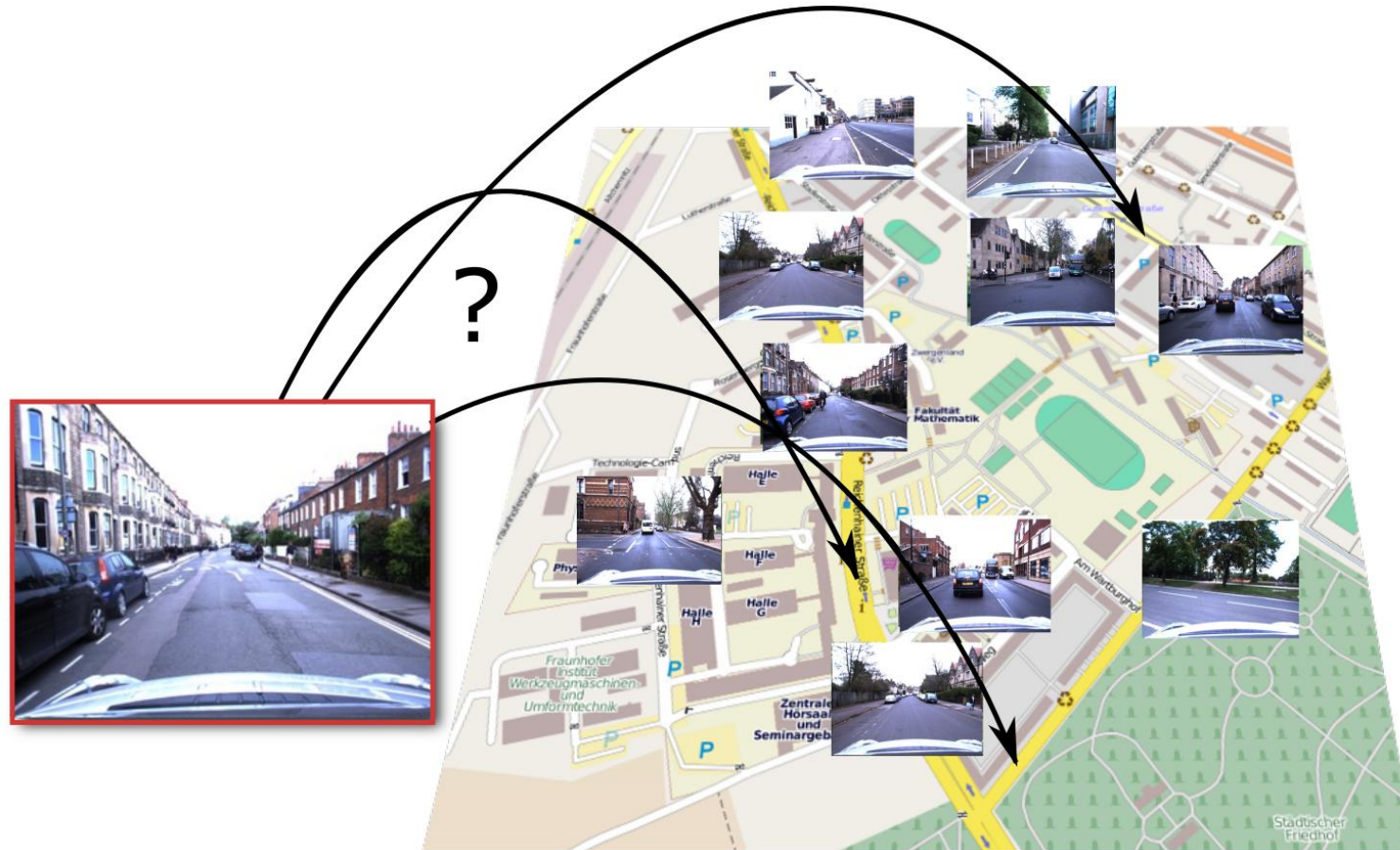
The Nordland dataset – a 3000 km journey across all four seasons



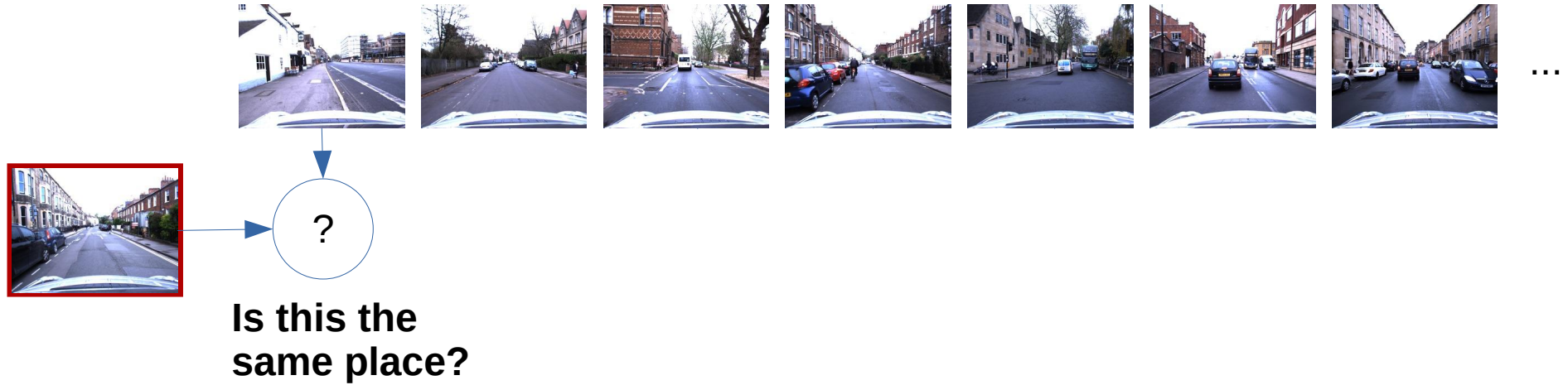
Sünderhauf, N., Neubert, P. & Protzel, P. (2013) Are We There Yet? Challenging SeqSLAM on a 3000 km Journey Across All Four Seasons. In Proc. of Workshop on Long-Term Autonomy at Int. Conf. on Rob. a. Autom. (ICRA)

<http://nrkbeta.no/2013/01/15/nordlandsbanen-minute-by-minute-season-by-season/>

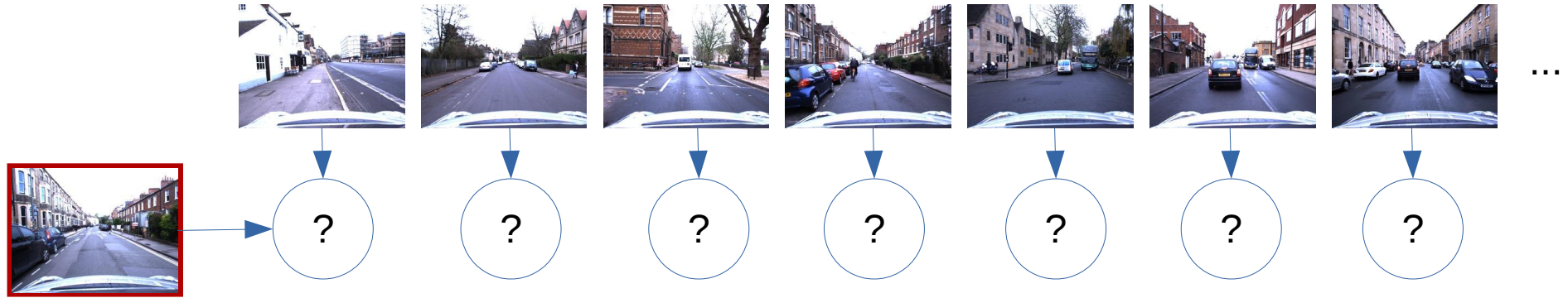
Teaser application 2: Place recognition in changing environments



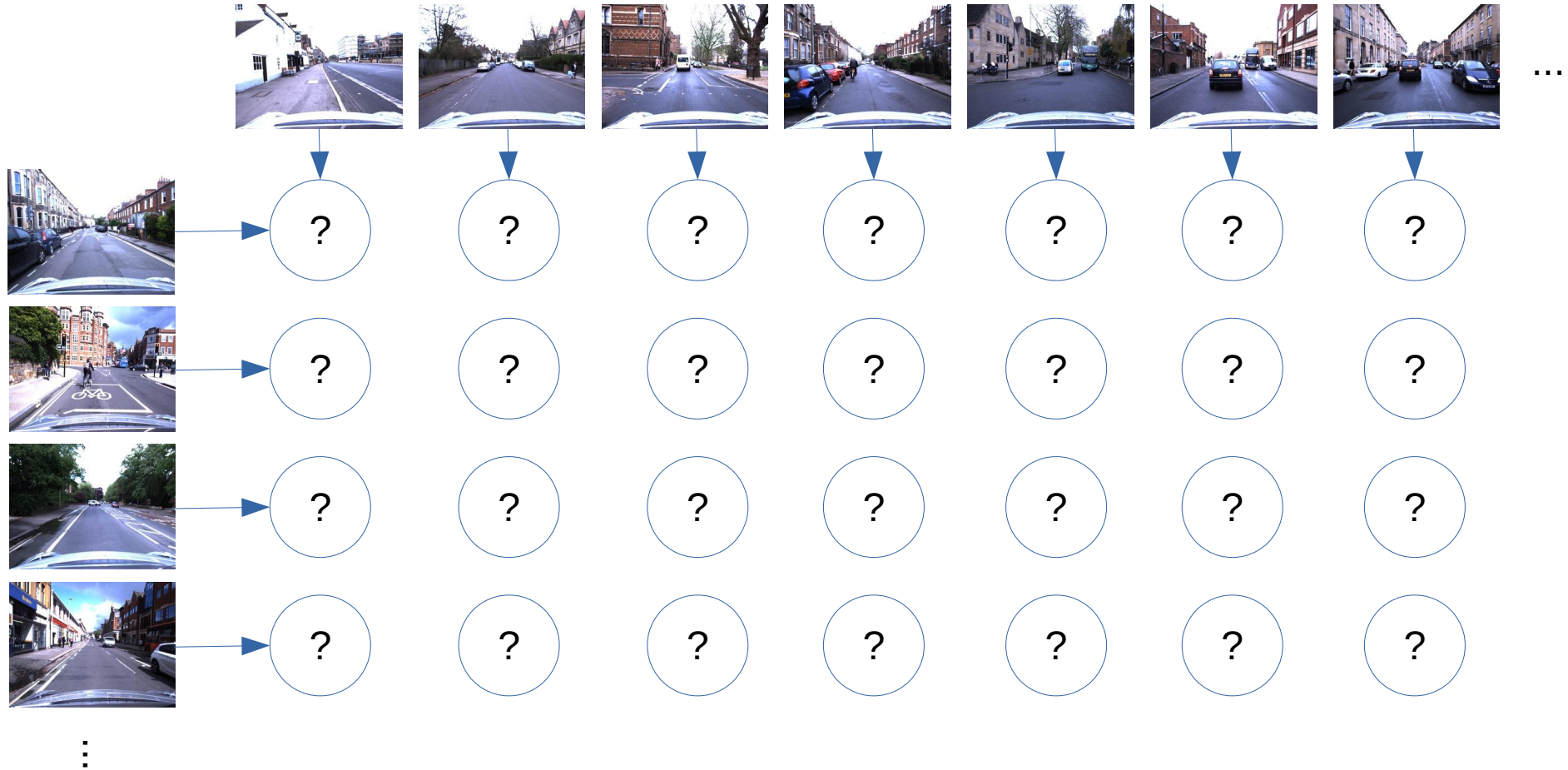
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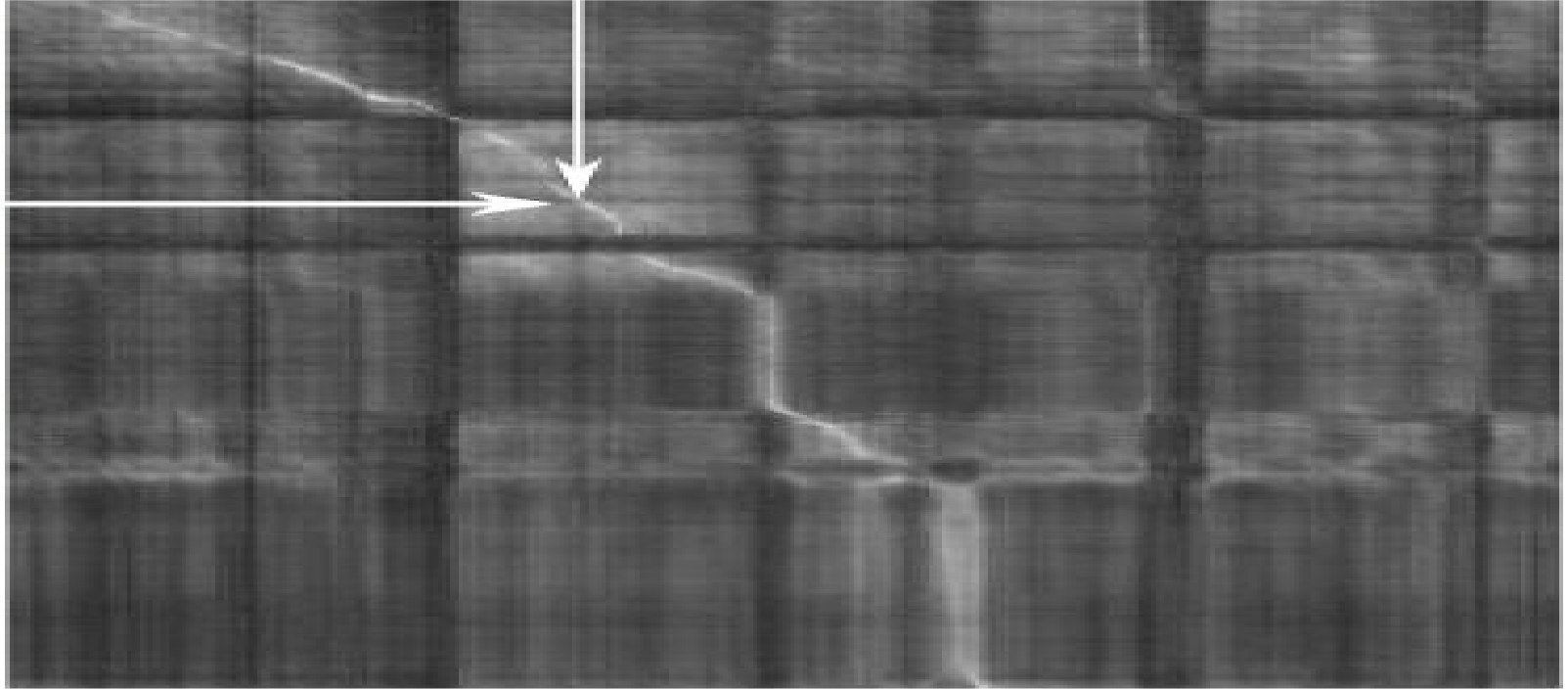
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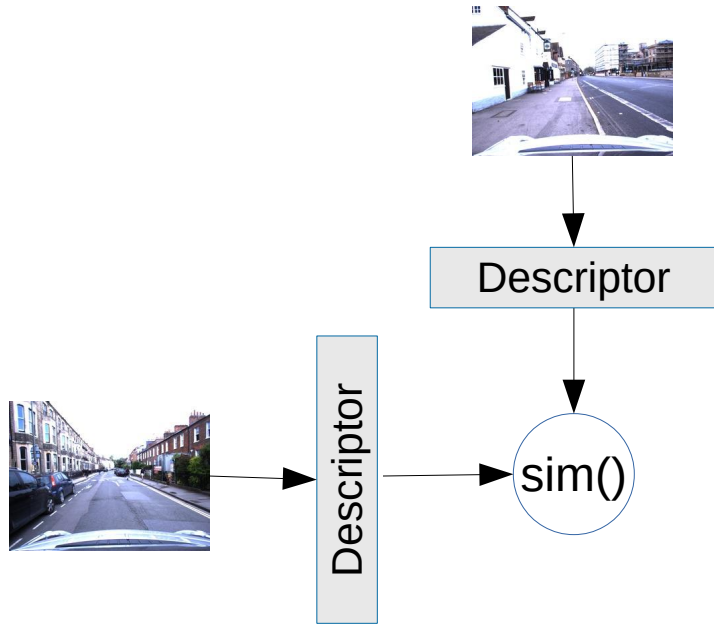
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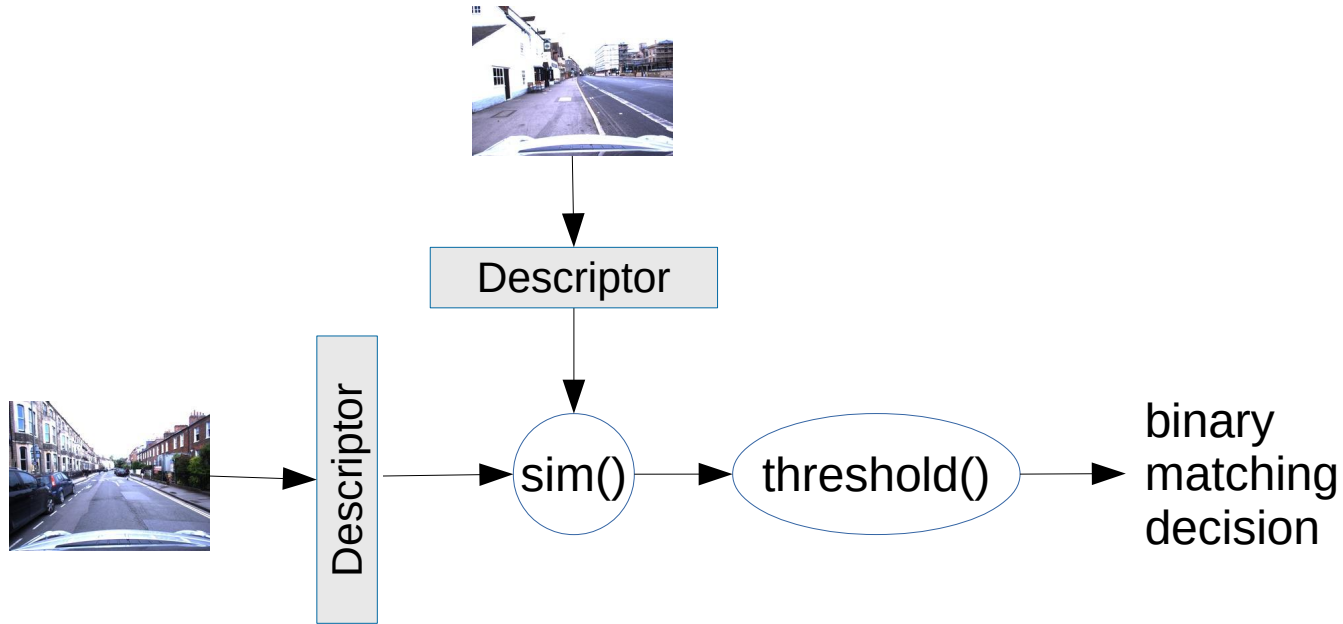
A simple approach: Pairwise descriptor matching



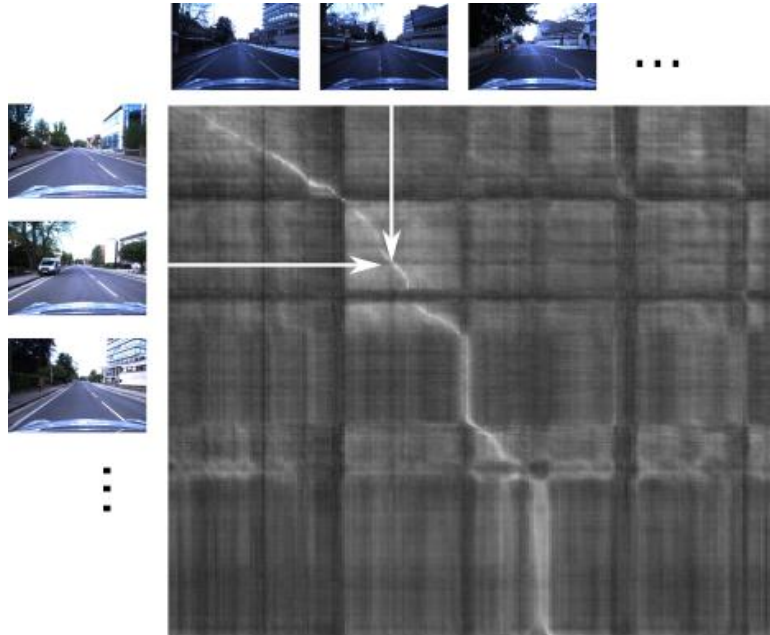
A simple approach: Pairwise descriptor matching



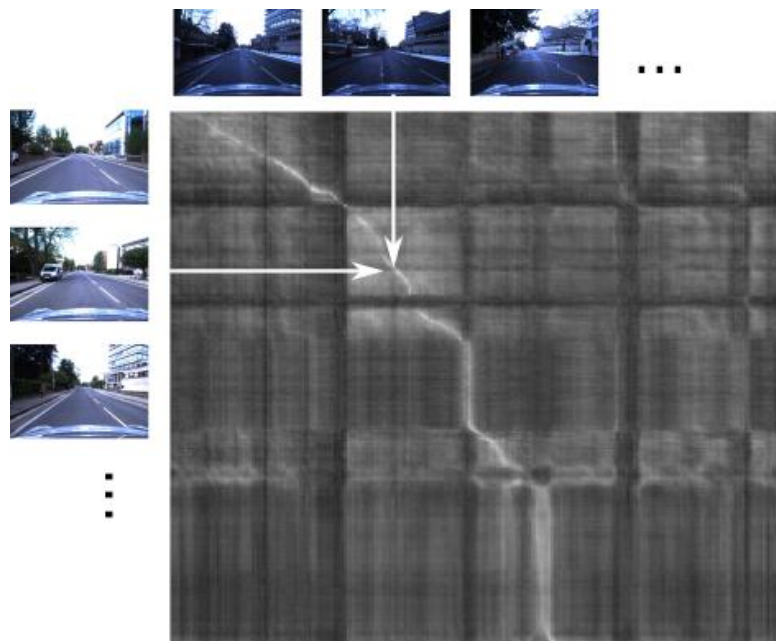
A simple approach: Pairwise descriptor matching



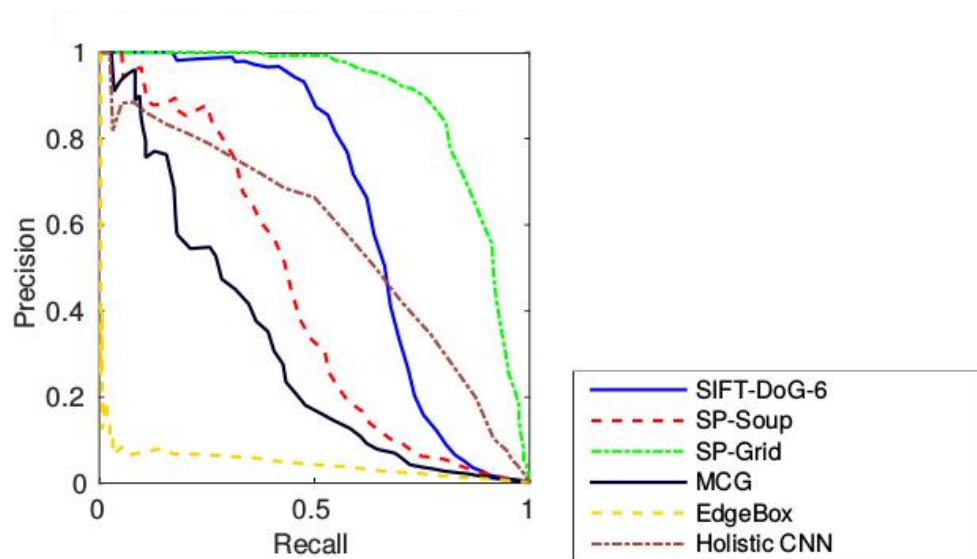
A simple approach: Pairwise descriptor matching



A simple approach: Pairwise descriptor matching



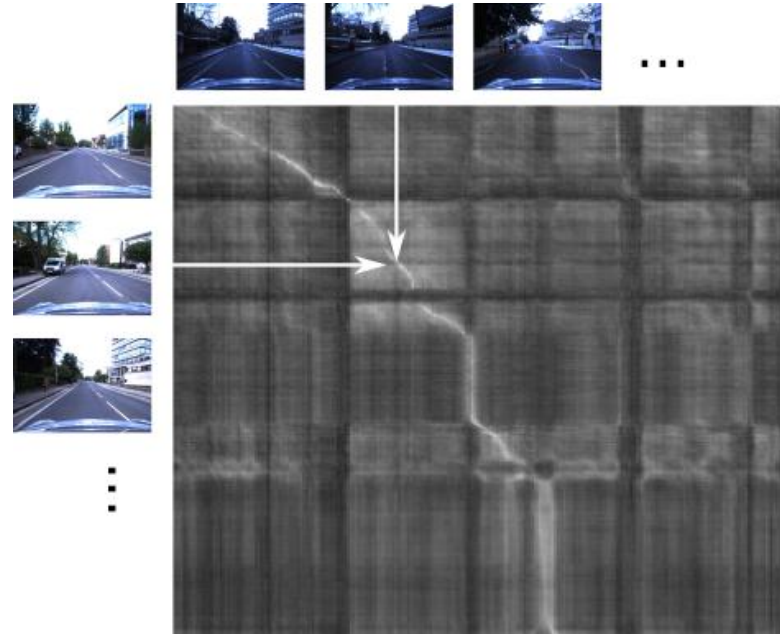
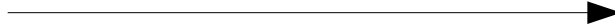
Evaluation based on
Precision-Recall curves



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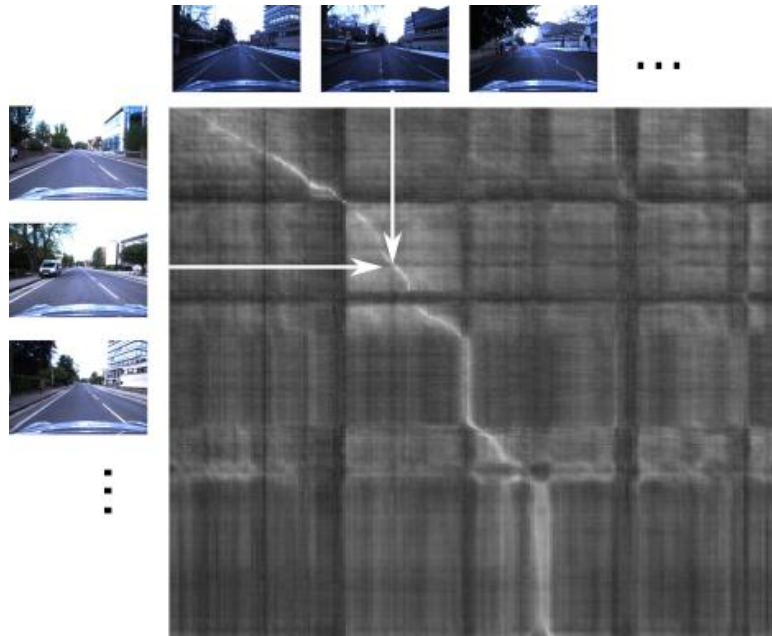


Deep
Neural
Network

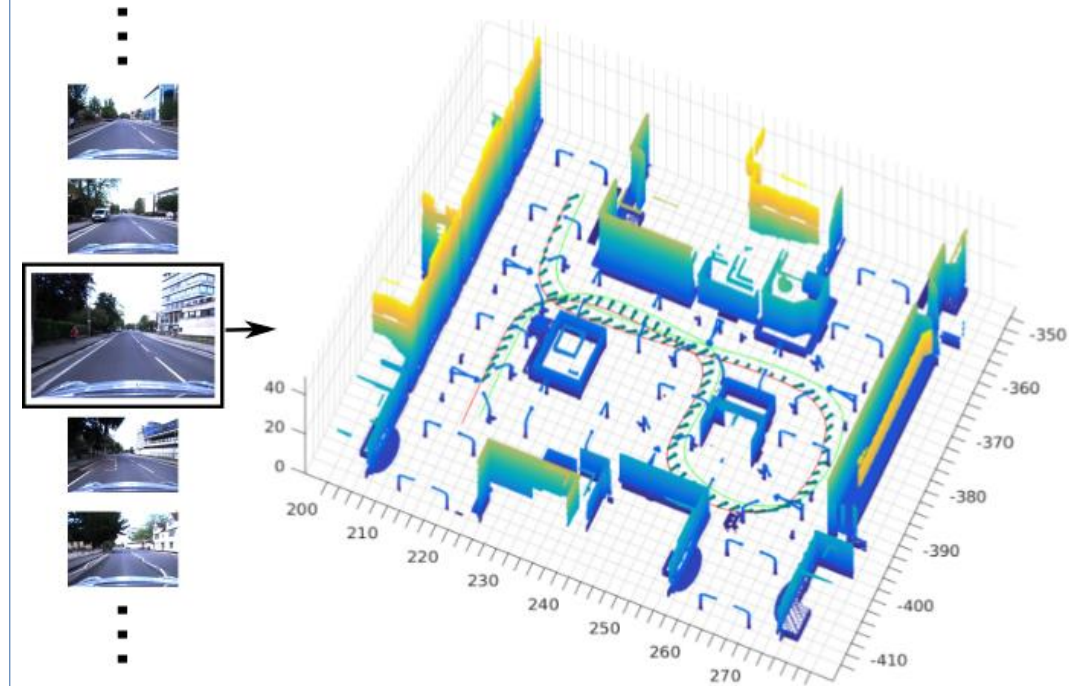

$$\begin{pmatrix} 1.0 \\ 3.9 \\ -0.5 \\ \cdot \\ \cdot \\ 2.9 \\ -6.0 \\ 9.8 \end{pmatrix}$$


Approaches to place recognition

Pairwise image comparison

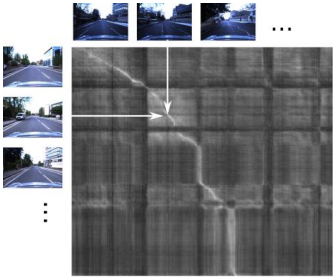


Metric SLAM

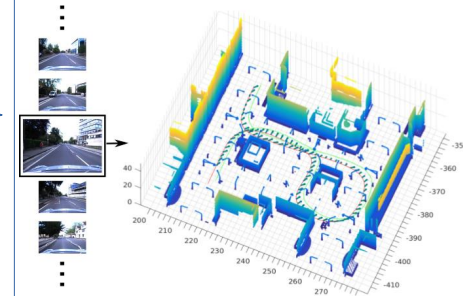


Approaches to place recognition

Pairwise comparison



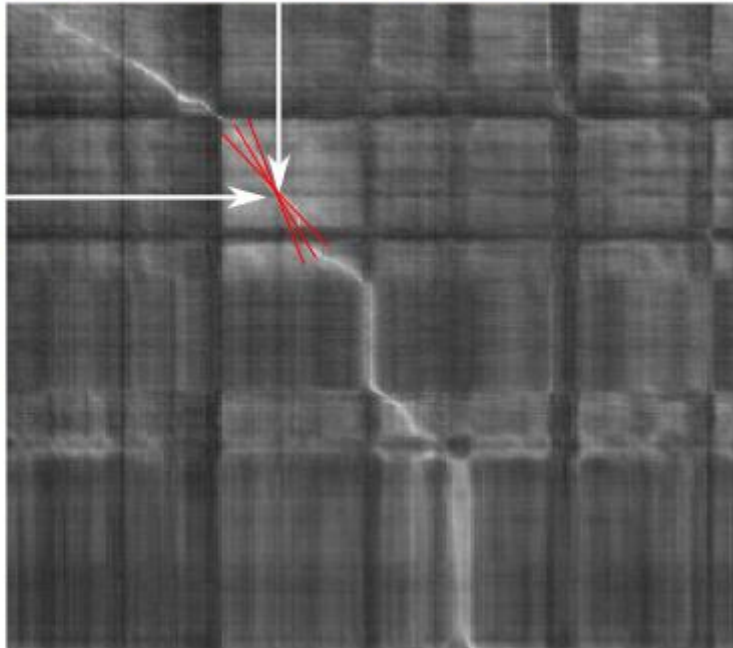
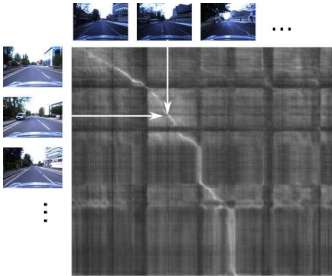
Metric SLAM



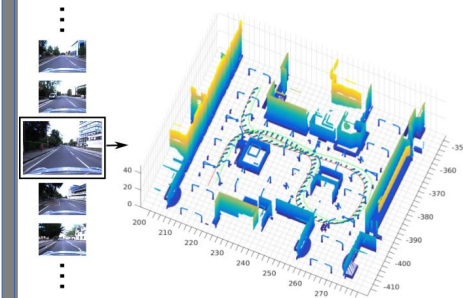
Approaches to place recognition

In between, e.g. SeqSLAM

Pairwise
independently



Metric SLAM



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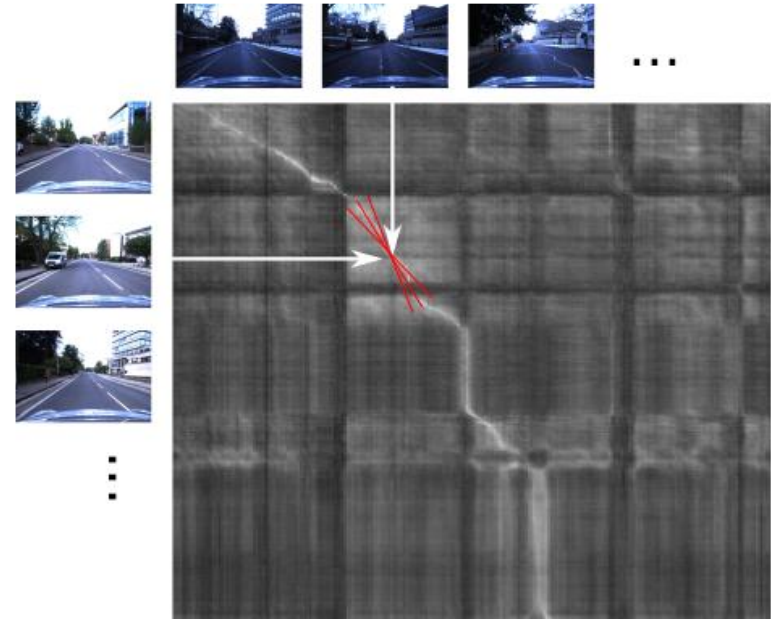


Deep
Neural
Network

$\begin{pmatrix} 1.0 \\ 3.9 \\ -0.5 \\ \vdots \\ \vdots \\ 2.9 \\ -6.0 \\ 9.8 \end{pmatrix}$

This is where the
hyperdimensional
magic happens

$$Y_i = \bigoplus_{k=0}^d (X_{i-k} \otimes P_{-k})$$



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Simplified SeqSLAM core:

input: distance matrix D

for each summer image idx j in S

for each winter image idx i in W

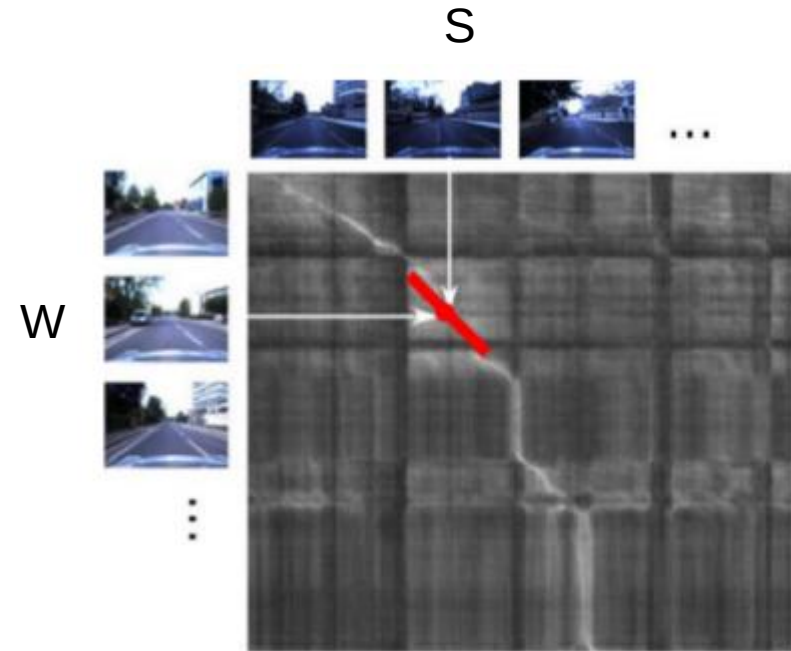
$\text{accDist} = 0$

for $k=-d:1:d$

$\text{accDist} = \text{accDist} + D(i+k, j+k)$

$R(i,j) = \text{accDist} / (2*d+1)$

output: resulting distance matrix R



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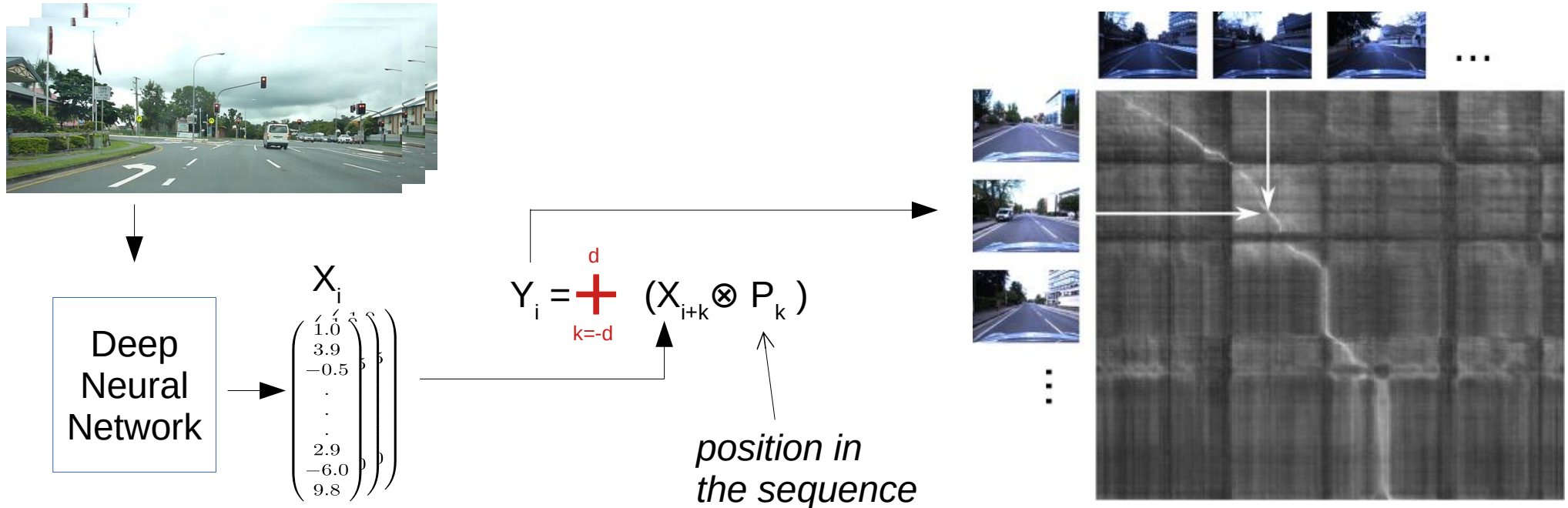
VSA approach

- Replace each image vector with a vector that represents the whole **sequence**
- Use this vector for the direct **pairwise** comparison

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VSA approach

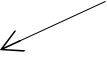
- Replace each image vector with a vector that represents the whole **sequence**
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$$Y_i = \bigoplus_{k=-d}^d (X_{i+k} \otimes P_k)$$

*position in
the sequence*



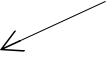
Why does this work?

e.g. comparing two 2-element sequences $A = (X_{a_1} X_{a_2})$
 $B = (X_{b_1} X_{b_2})$

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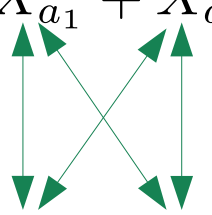
position in the sequence



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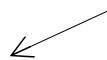
Without binding to position

$$Y_A = X_{a_1} + X_{a_2}$$

$$Y_B = X_{b_1} + X_{b_2}$$

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position in the sequence



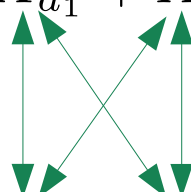
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Without binding to position

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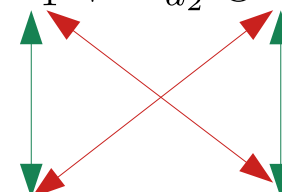
$$Y_B = X_{b_1} + X_{b_2}$$



With binding to position

$$Y_A = X_{a_1} \otimes P_1 + X_{a_2} \otimes P_2$$

$$Y_B = X_{b_1} \otimes P_1 + X_{b_2} \otimes P_2$$



Hands-on