

A Completion Network for Reconstruction from Compressed Acquisition

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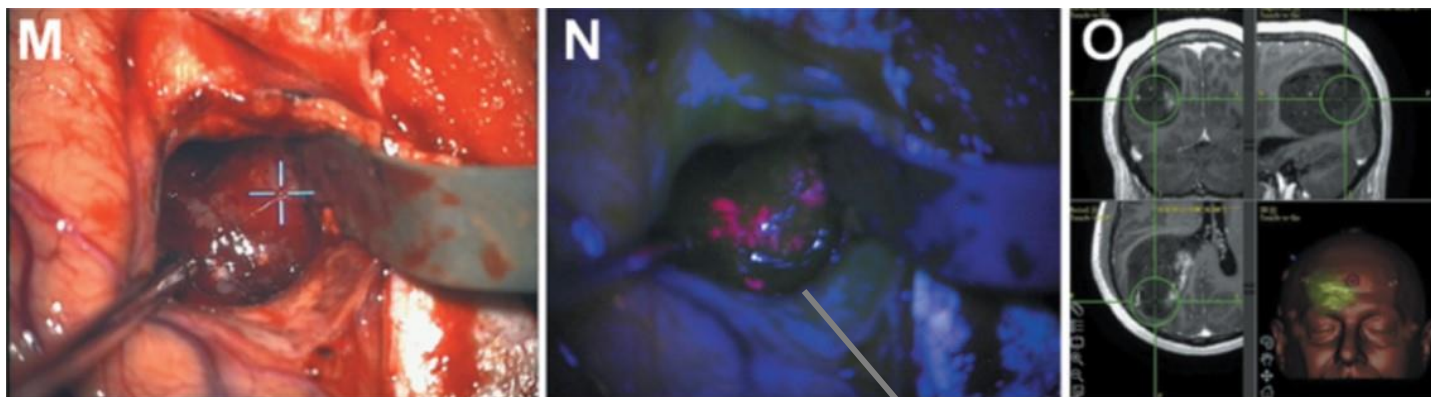


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➤ Fluorescence-guided neurosurgery

- ❖ Protoporphyrin IX (PpIX) fluorescence

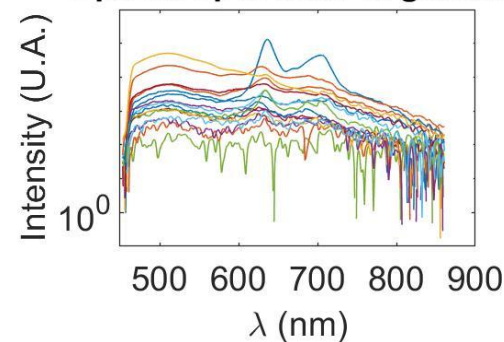
P. Valdés *et al.*, *J. Neurosurgery*, **123**(3), 2015



➤ Full spectrum acquisition

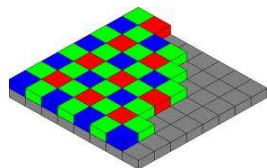
- ❖ Better **detection** of tumor **margins**
[P. Valdés *et al.*, *JNS*, **123**(3), 2015]
[P. Leclerc *et al.*, *Sci Reports* **10**, 2020]

Optical spectrum of gliomas



Array

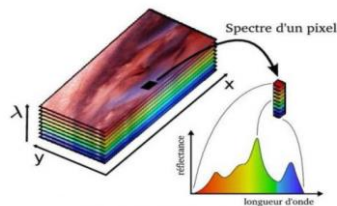
Colour



Multi-spectral



Hyper-spectral



Point

Spectrometer



Spatial resolution

yes

yes

yes

Spatial resolution

no

Number of spectral channels

3

2—10

10—100

Spectral channels

100—2,000

Cost

~€1k

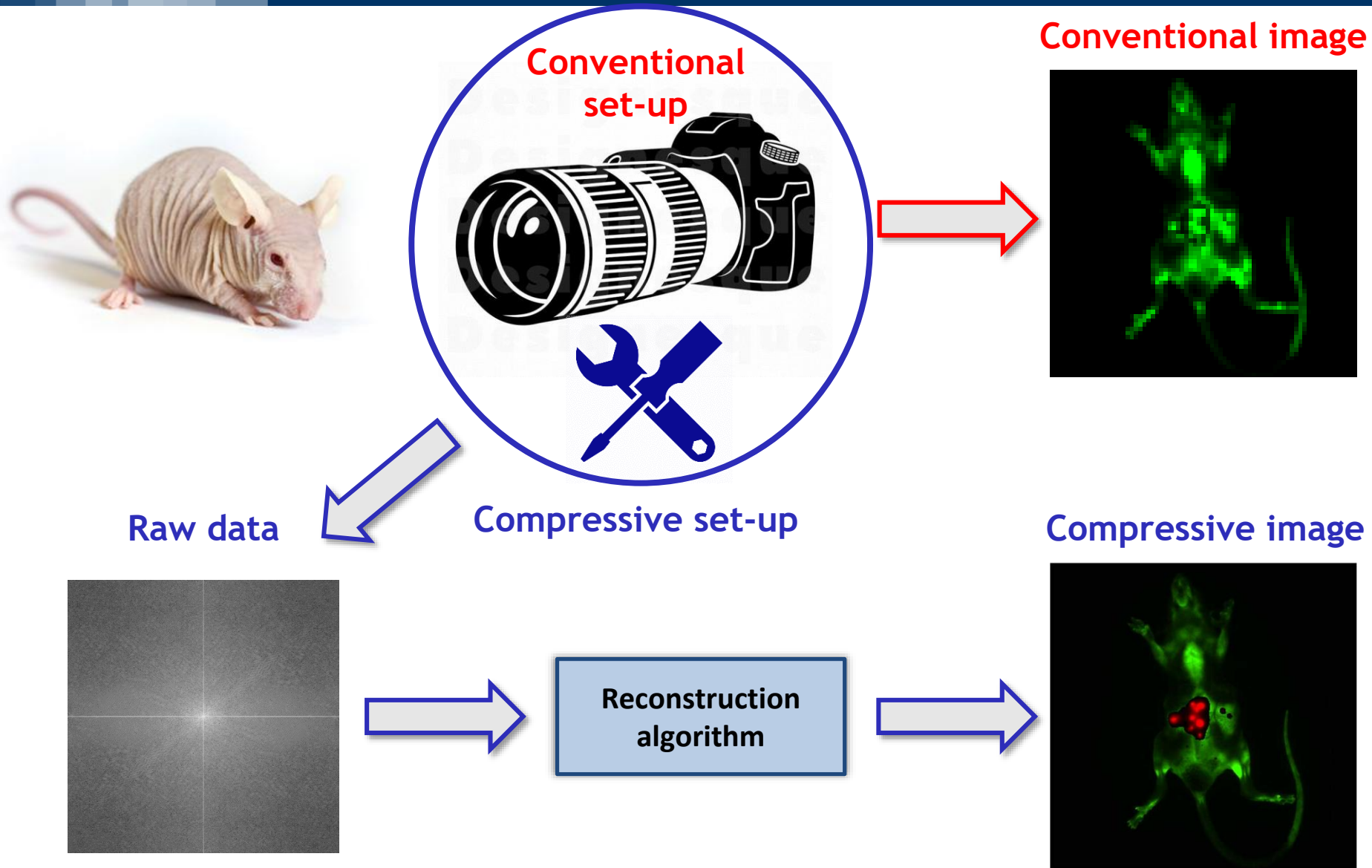
~€10k

~€100k

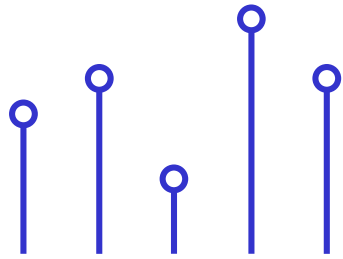
Cost

~€1k

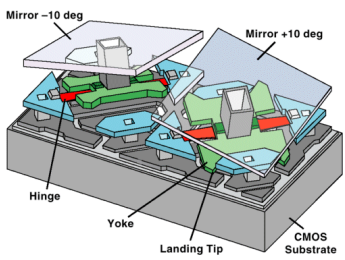
Need for **low-cost array** with **high spectral resolution**



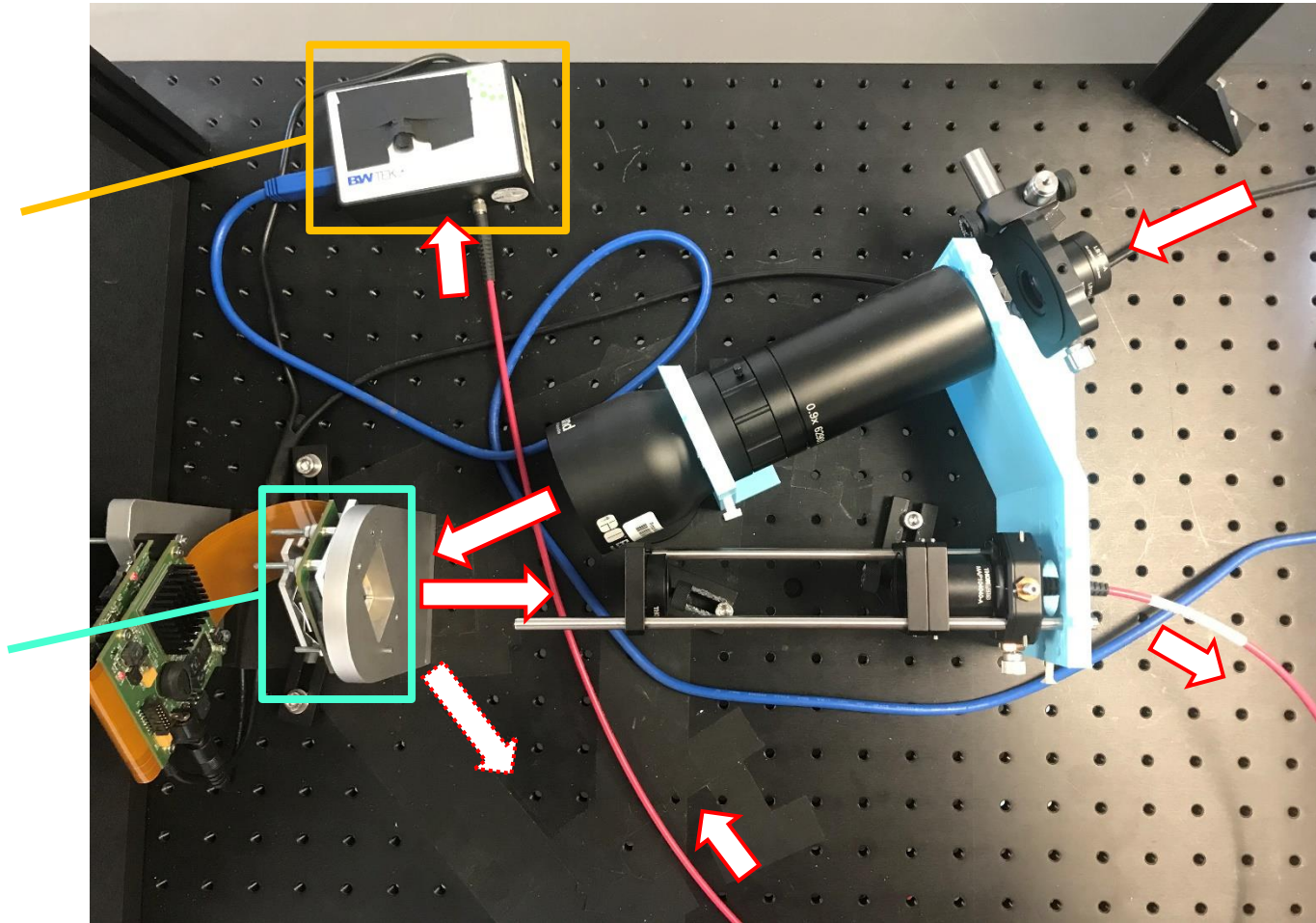
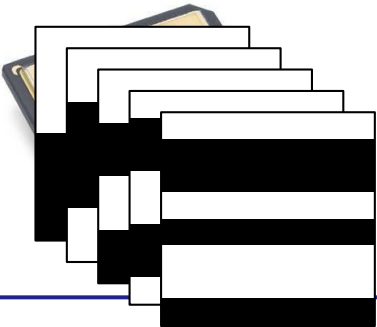
COMPRESSIVE (SINGLE-PIXEL) CAMERA

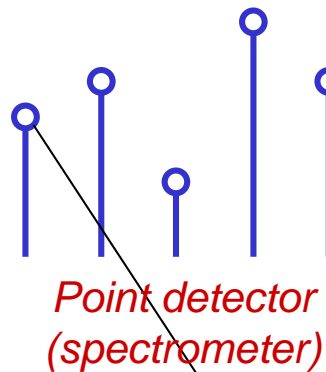


Point detector (spectrometer)



Spatial light modulator

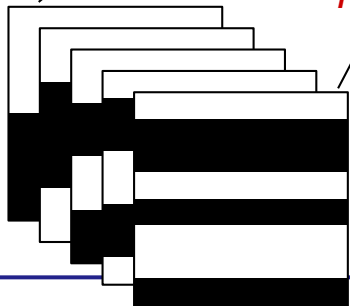




$$\mathbf{m} = [m_1, \dots, m_M]^T \in \mathbb{R}^M$$

$$\mathbf{P}_1 = [\mathbf{p}_1^T, \dots, \mathbf{p}_M^T]^T \in \mathbb{R}^{M \times N}$$

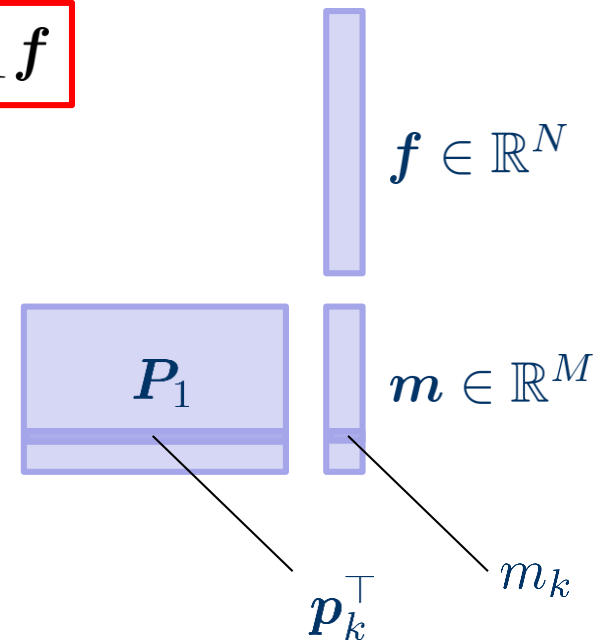
Spatial
light
modulator



➤ Linear model

$$\mathbf{m} = \mathbf{P}_1 \mathbf{f}$$

$$M \ll N$$



➤ Challenge

- ❖ A small M limits the acquisition time
- ❖ A small M limits the image resolution too!

1. **Experiment design:** How to choose the patterns (codes) P ?
Not addressed here! We choose

$$m = P_1 f$$

orthogonal basis $\rightarrow P = \begin{bmatrix} P_1 \\ P_2 \end{bmatrix}$ \leftarrow **acquired** patterns
 \leftarrow **missing** patterns

2. **Reconstruction:** How to recover the image f from m ?

❖ Constrained optimization

$$\min_f \mathcal{R}(f) \quad \text{such that} \quad m = P_1 f.$$

- Least squares: fast but low resolution [Rousset et. al, IEEE TCI, 2017]
- Total variation: higher resolution but time consuming [Duarte et. al, IEEE SPM, 2009]

❖ CNN: Learn a nonlinear reconstructor [Higham et. al, Sci. Rep., 2018]

$$f^* = \mathcal{H}_\theta(m),$$

➤ The least-squares problem

$$\min_f \|\mathbf{f}\|_2^2 \quad \text{such that} \quad \mathbf{m} = \mathbf{P}_1 \mathbf{f}.$$

...has the closed-form solution

$$\mathbf{f}^* = \mathbf{P}_1^\top \mathbf{m}$$

... equivalent to

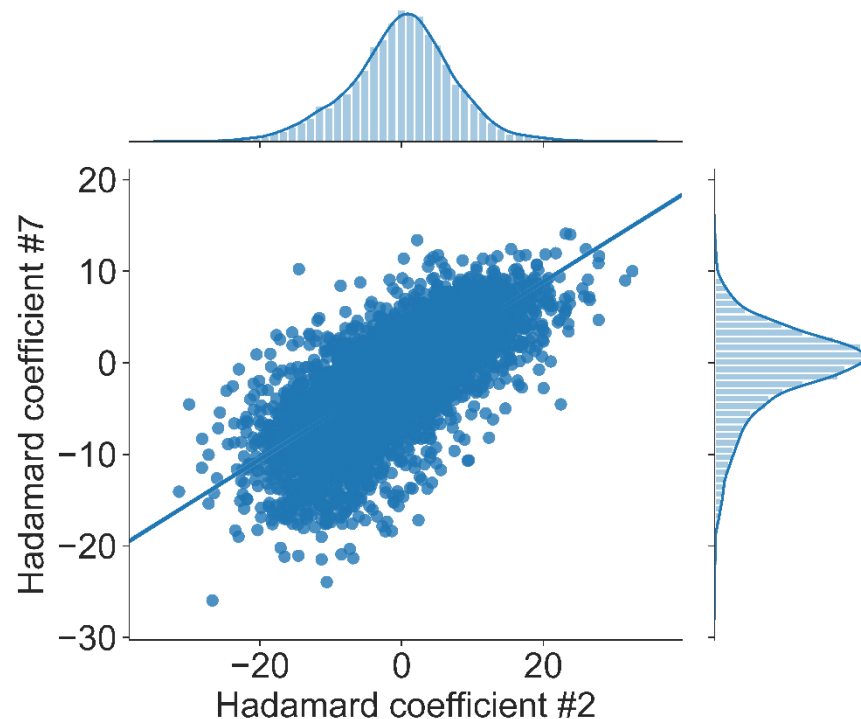
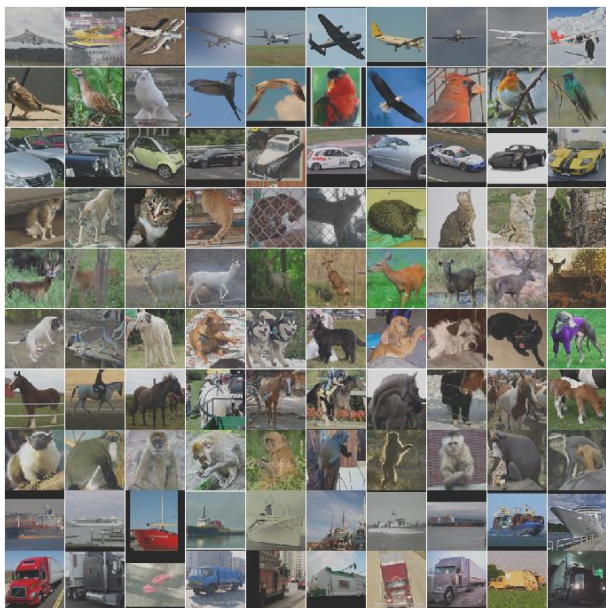
$$\mathbf{f}^* = \mathbf{P}^\top \mathbf{y}^*, \quad \text{with } \mathbf{y}^* = \begin{bmatrix} \mathbf{m} \\ \mathbf{0} \end{bmatrix} \in \mathbb{R}^N$$



What about completing the missing measurements by relevant values?

➤ How to complete?

STL-10 dataset



➔ Exploiting the correlation between the measured coefficients

Completion approach

$$\mathbf{f}^* = \mathbf{P}^\top \mathbf{y}^*, \quad \text{with } \mathbf{y}^* = \begin{bmatrix} m \\ \mathbf{y}_2^* \end{bmatrix},$$

with

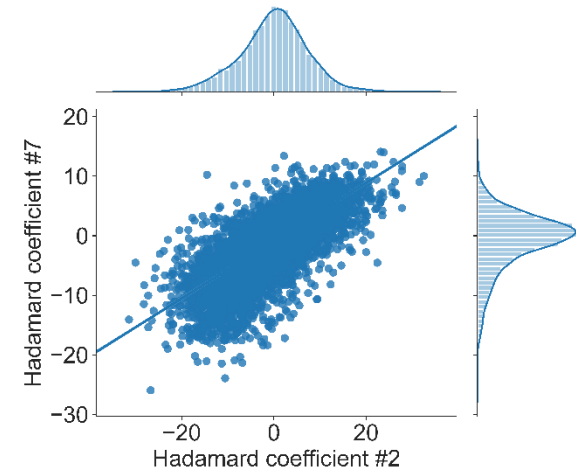
$$\mathbf{y}_2^*(m) = \mathbb{E}(\mathbf{y}_2 \mid \mathbf{y}_1 = m)$$

➤ **Under Gaussian assumptions**

$$\mathbf{y}_2^*(m) = \boldsymbol{\mu}_2 + \boldsymbol{\Sigma}_{21} \boldsymbol{\Sigma}_1^{-1} (m - \boldsymbol{\mu}_1)$$

Covariance between
measured and missing

Covariance of
measured

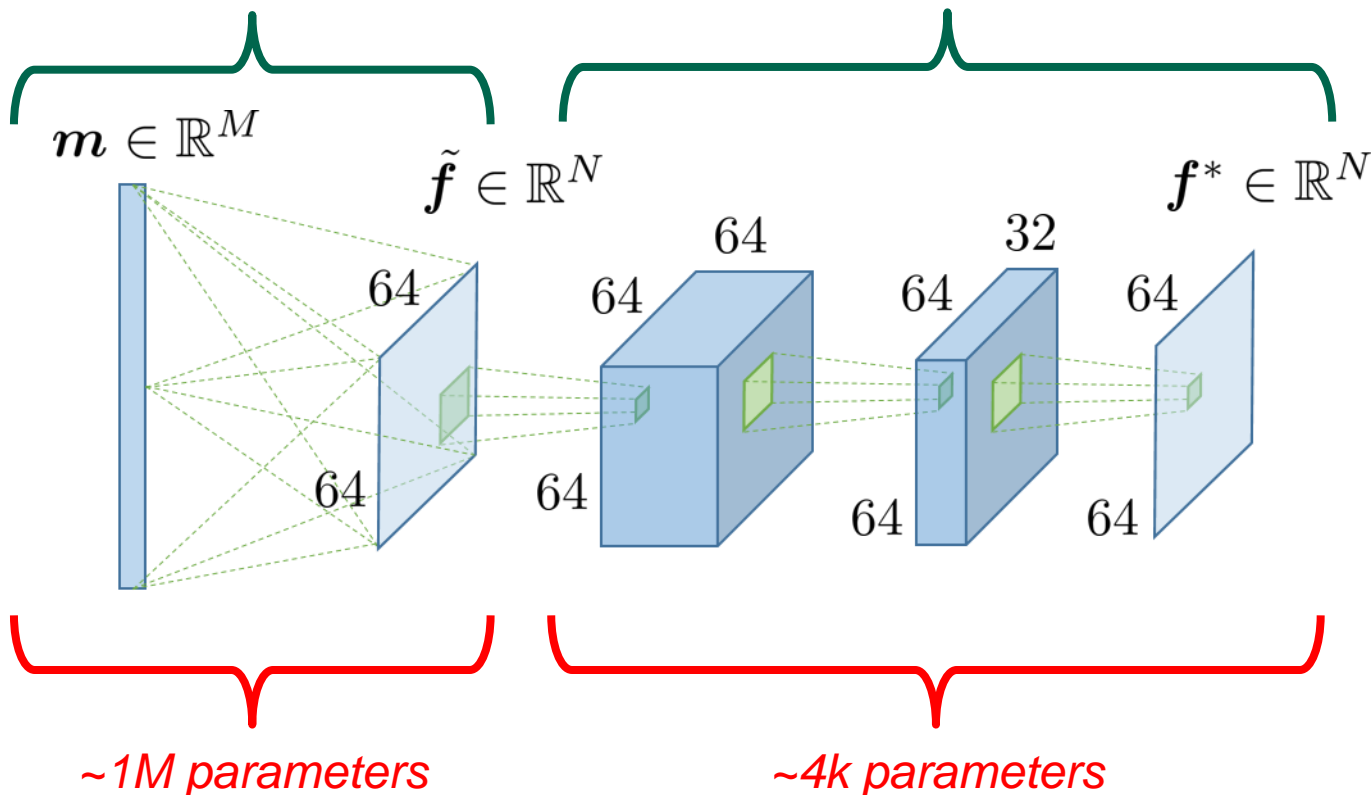


➤ **No assumption: This is the best linear solution!**

➤ CNN architecture

Fully-connected layer (FCL)

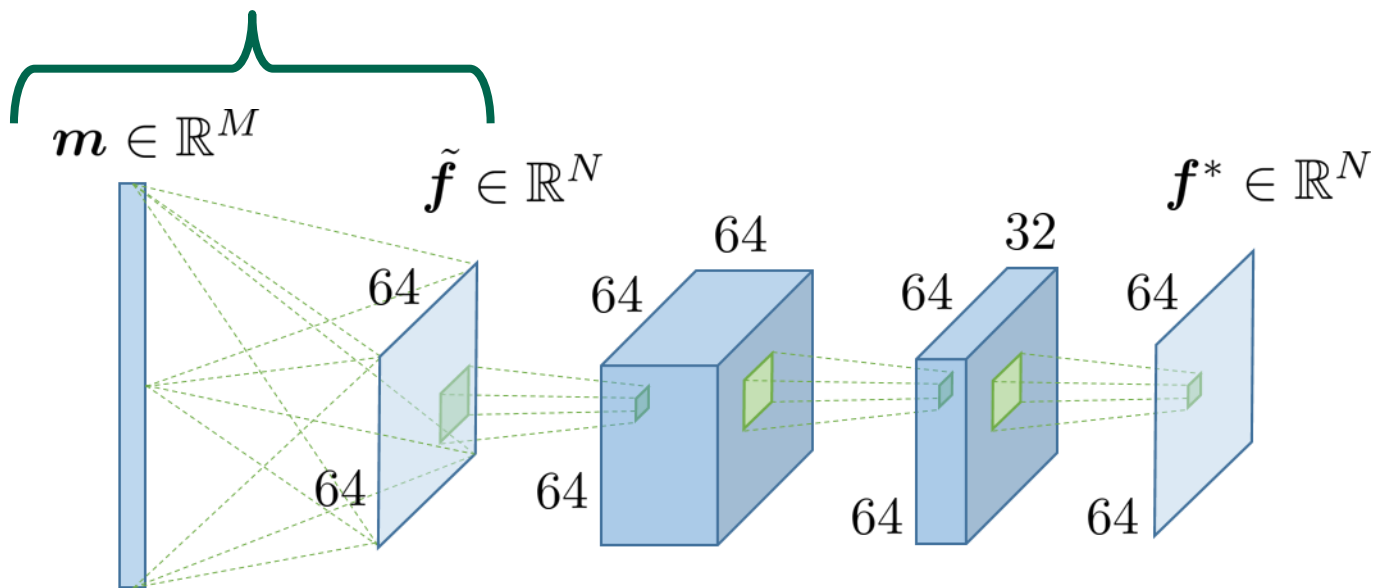
Convolutional layers



[Higham *et. al*,
Sci. Rep., 2018]

➤ CNN architecture

Fully-connected layer (FCL)

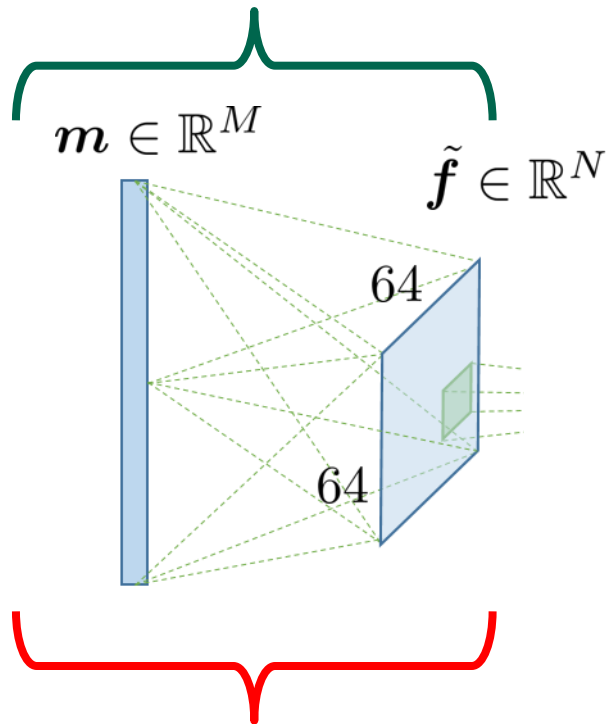


Gaussian completion recon

*Learn to compensate for
the non Gaussianities*

➤ CNN architecture

Fully-connected layer (FCL)



measurement-to-image
mapping

➤ Choices for the FCL

- ❖ Free [Higham et. al, Sci. Rep., 2018]

$$\tilde{\mathbf{f}} = \mathcal{H}_{\theta_1}(\mathbf{m})$$

- ❖ Pseudo inverse [Jin et. al, IEEE TIP, 2017, Ravishankar et. al, Proc. IEEE, 2020]

$$\tilde{\mathbf{f}} = \mathbf{P}_1^\top \mathbf{m}$$

- ❖ Bayesian completion

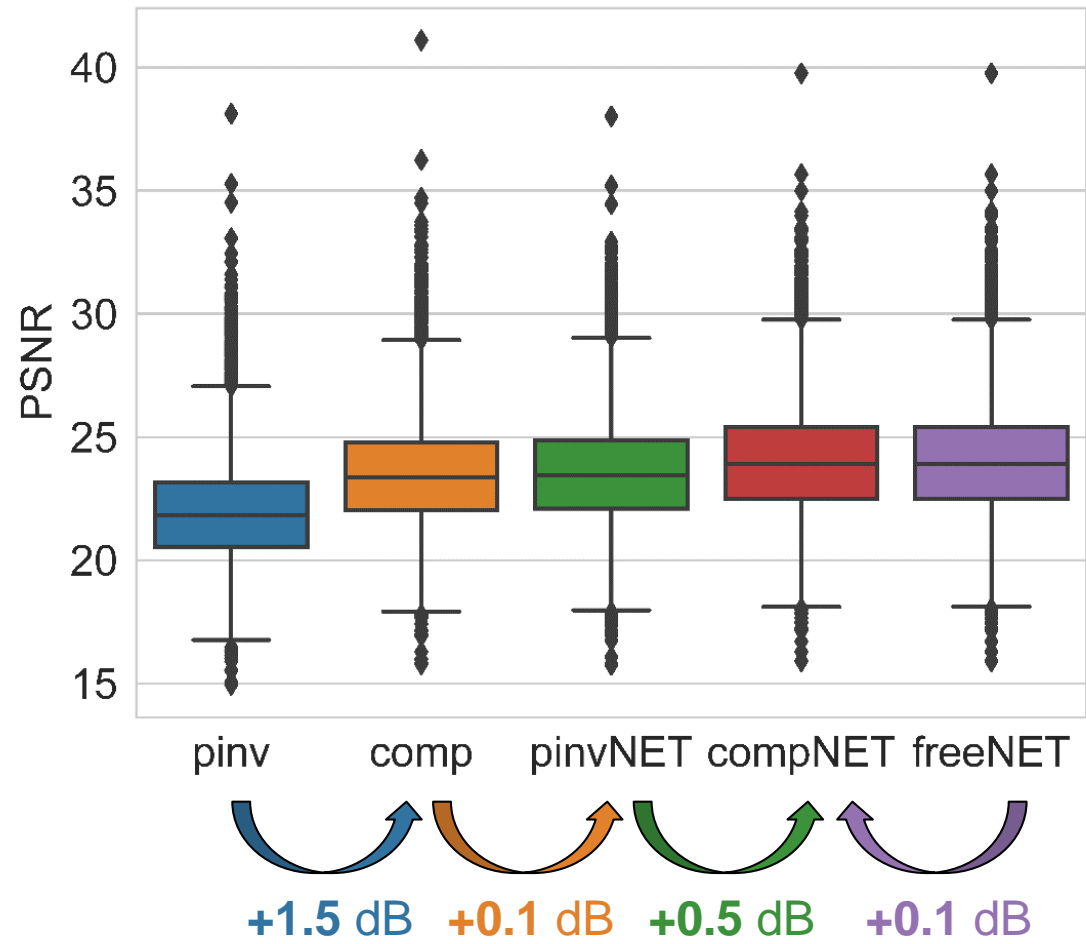
$$\tilde{\mathbf{f}} = \mathbf{P}^\top \mathbf{y}^*, \quad \text{with } \mathbf{y} = \begin{bmatrix} \mathbf{m} \\ \mathbf{y}_2^* \end{bmatrix}$$

➤ 3 network variants

- ❖ freeNet: (~1M parameters)
- ❖ pinvNet: (~4k parameters)
- ❖ compNet: (~4k parameters)

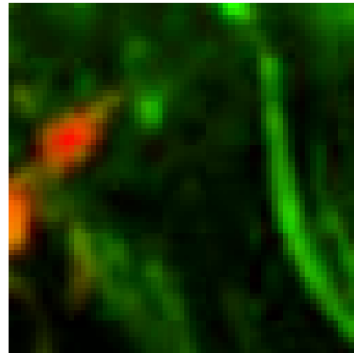
➤ **STL-10 (Training using ~100k images, test using 8k images)**

pinv: 22.0 ± 2.2 dB
comp: 23.5 ± 2.2 dB
pinvNET: 23.6 ± 2.2 dB
compNET: 24.1 ± 2.3 dB
freeNET: 24.0 ± 2.2 dB



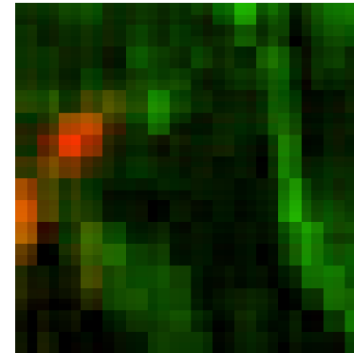
➤ **Fluorescence microscopy images (not from STL-I0!)**

(a) Ground-Truth



f^{true}

(b) Pseudo Inverse

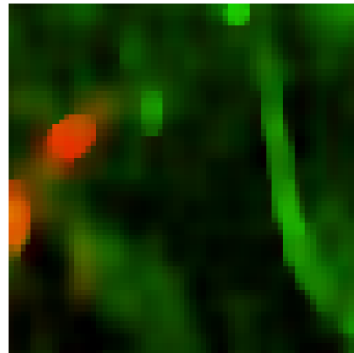


$$f^* = \arg \min_f \|f\|_2^2$$

s. t. $m = Pf$

red: 27.15 dB
green: 24.27 dB

(c) Total Variation

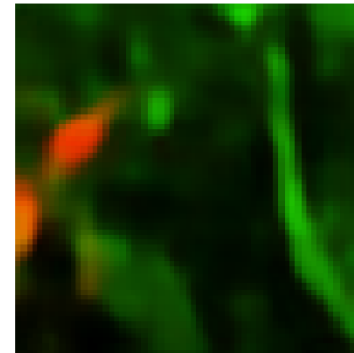


$$f^* = \arg \min_f \|\nabla f\|_1$$

s. t. $m = Pf$

red: PINV + 3.52 dB
green: PINV + 2.41dB

(d) compNET



$$f^* = \mathcal{H}_{\theta^*}(m)$$

red: TV + 0.8 dB
green: TV + 1.16 dB

➤ Conclusions

- ❖ Reconstruction as completion
- ❖ Simple **linear reconstruction** scheme based on **Bayesian completion**
- ❖ Simple **nonlinear convolutional network**
- ❖ **Code**
 - Bayesian completion (MATLAB): <https://github.com/nducros/SPIRIT>
 - Convolutional Network (Python): soon...

➤ Perspectives

- ❖ Noise
- ❖ Experimental data
- ❖ Video imaging
 - Much higher compression rate
 - Still challenging...

Don't miss the talk by Antonio Lorente Mur Tomorrow morning!

Thanks for watching!