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AAIC > 23

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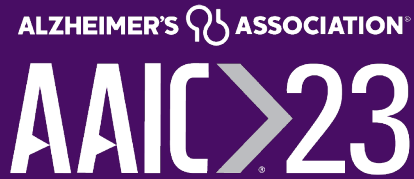
JULY 16-20 > AMSTERDAM, NETHERLANDS, AND ONLINE

ISTAART Neuroimaging PIA THE BASICS OF NEUROIMAGING SEMINAR SERIES



ISTAART Neuroimaging PIA

The Basics of Neuroimaging Series



BASICS OF NEUROIMAGING

Positron Emission Tomography (PET)

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@tobeybetthouser

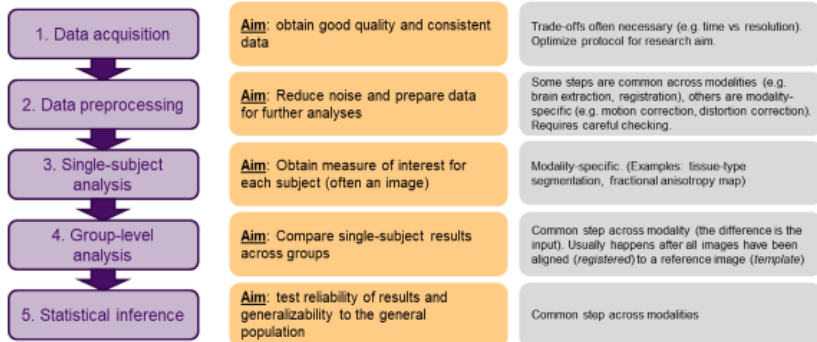


tjbetthouser@medicine.wisc.edu

Basics of Neuroimaging: Data Structure and Formats

Dr. Ludovca Griffanti

ALZHEIMER'S ASSOCIATION AAIC>23 NEUROIMAGING DATA ANALYSIS: A GENERIC BLUEPRINT



Basics of Neuroimaging: Structural MRI

Dr. David Cash

ALZHEIMER'S ASSOCIATION AAIC>23 MRI SEQUENCES AND WEIGHTINGS



Available On-Demand soon at: <https://training.alz.org/research-webinars>

By the end of this session, you should be able to:

- Understand the differences between PET imaging and other modalities
- Understand how PET imaging data is collected and image are created
- Perform basic PET imaging processing and quantification for tracers commonly used in ADRD

Basics of PET Imaging

- How is PET different from other techniques?
- What is a PET tracer?
- How do we get an image?
- How do we quantify PET?

PET Image Processing

- MR-guided
- PET-Only
- Other Considerations

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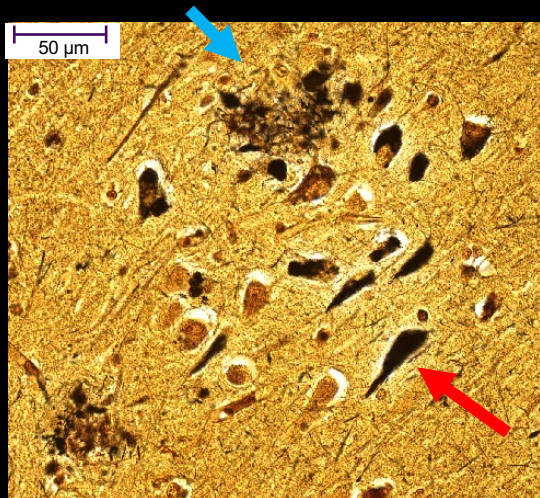
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Introduction – Basics of PET

- **How is PET different from other imaging modalities?**
- **What is a PET tracer?**
- How do we get a PET image?
- How do we quantify PET?

How is PET different from other techniques?

Microscopy



~μm resolution

Beta-amyloid plaque

Neurofibrillary tau tangle

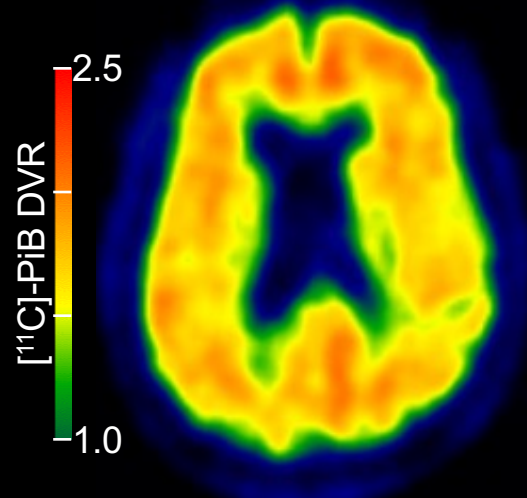
Structural MRI



~1 mm resolution

Brain Volume and Anatomy

PET



~4-6 mm resolution

Biology/Physiology

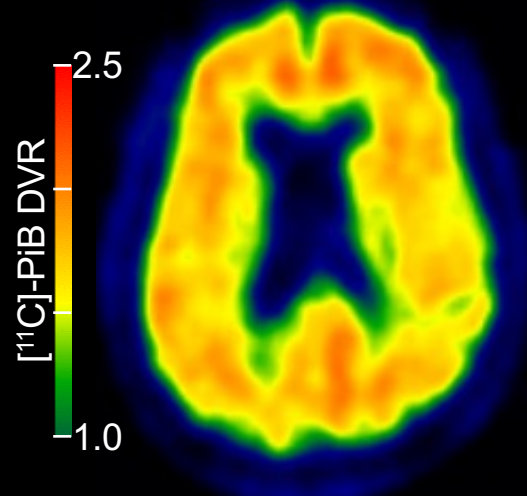
How is PET different from other techniques?

Some things PET can measure:

- Binding potential (proportional to receptor density)
- Rate Constants (e.g., influx/efflux from/to plasma and tissue, binding to and dissociation from target)
- Tissue Perfusion and Relative Perfusion
- Receptor Occupancy
- Metabolic Rate (FDG)

Low Spatial Resolution, High Molecular Specificity

PET

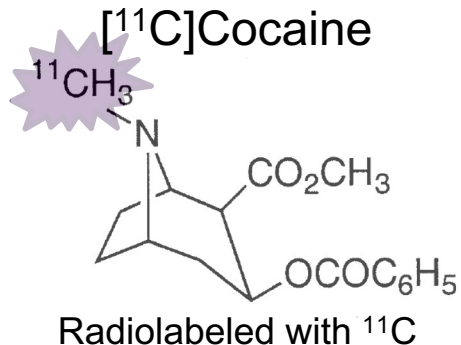
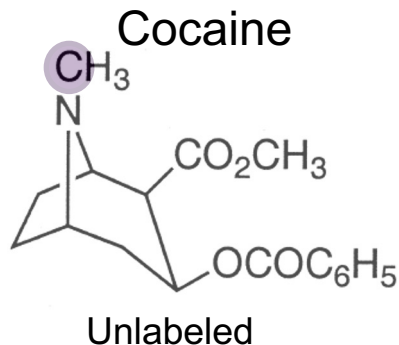


~4-6 mm resolution
Binding Potential

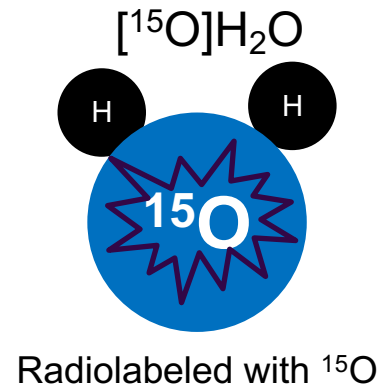
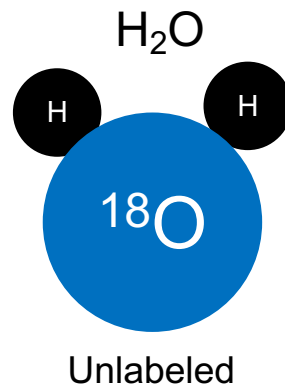
What is a PET tracer?

- A molecule we want to image with a positron emitting isotope attached (i.e., radiolabeled)

Cocaine as a PET tracer of DAT

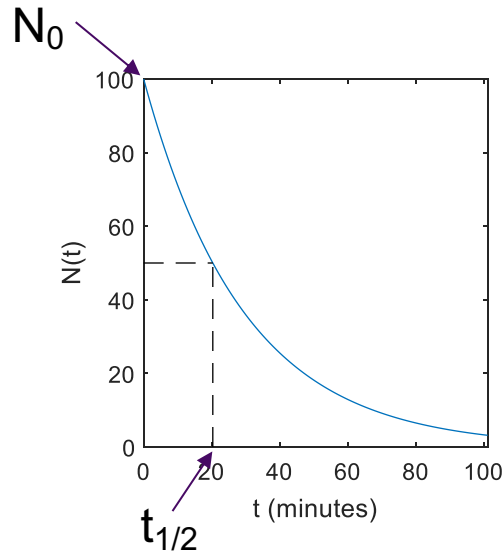


Water as a PET tracer of Perfusion



What is a PET tracer?

- A molecule we want to image with a positron emitting isotope attached (i.e., radiolabeled)
- A radioactive isotope is a form of an element (e.g., carbon) with an unstable nucleus that undergoes radioactive decay



$$N(t) = N_0 e^{-\frac{\ln(2)t}{t_{1/2}}}$$

$t_{1/2}$ = half life

What is a PET tracer?

- A molecule we want to image with a positron emitting isotope attached (i.e., radiolabeled)
- A radioactive isotope is a form of an element (e.g., carbon) with an unstable nucleus that undergoes radioactive decay
- Match radioactive half-life to biological process we're trying to detect

PET Isotope	Half-life (minutes)
^{11}C	20.4
^{13}N	10.0
^{15}O	2.1
^{18}F	109.8

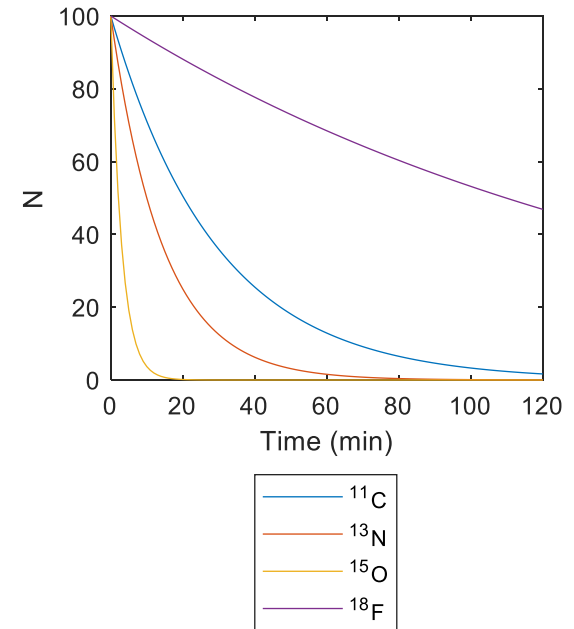


Table 5 Representative examples of radiotracers for CNS applications that have shown utility in humans

Targets	Carbon-11 labeled	Fluorine-18 labeled	Comments
β-Amyloid	[11C]PBB	[18F]flutemetamol [18F]florbetapir ([18F]AV-45) [18F]AZD 4694 [18F]FBI [18F]DDNP [18F]-SMBIB-W372 (F-18)-W372 [18F]florbetaben [18F]MK3328	
Tau		[18F]T807 (AV1451; Flortaucipir) [18F]GTP1 [18F]RO6958948 [18F]MK6240 [18F]PI-2620	Relative sensitivity to 3-repeat to 4-repeat tau isoforms remains to be confirmed.
Enzymes			
Aromatic amino acid decarboxylase (AADC)		6-[18F]-L-DOPA (FDOPA)	Used to assess dopamine synthesis capacity and storage; providing an indirect measure of functional integrity of the nigrostriatal dopaminergic pathway.
AChE	[11C]MP4A		
Aromatase	[11C]VOR		
FAAH	[11C]CURB		
MAO-A	[11C]flamine [11C]loglyline [11C]belixatone		
MAO-B	[11C]Deprenyl-d2		May be used as a marker of astrocytes
PDE4	[11C](R)-Rolipram		
PDE10A	[11C]DMA107 [11C]MP-10 [11C]Lu AB92686	[18F]MNI659	
Receptors			
Adenosine A1		[18F]CFFPX	
Adenosine A2A	[11C]SCH442416	[18F]MNI444	
GABAA	[11C]Flumazenil	[18F]flumazenil	
GABAA (alpha 5 preferring)	[11C]Ro15 4513		
CBI	[11C]MePPEP [11C]OMAR [11C]SD604	[18F]EMMEP-d2 [18F]MK-9470	
D1	[11C]NNC 112 [11C]SCH 23390		
D2/D3	[11C]Raclopride [11C]FLB 457 [11C]MPPA (agonist) [11C](+)-PHNO (agonist) [11C]NBA (agonist)	[18F]fallypride	
H1	[11C]Doxepin		
H3	[11C]GSK189254 [11C]GR 103545	[18F]FMH3	
5-HT1A	[carbonyl-11C]WAY [carbonyl-11C]DWAY	[18F]CWAY [18F]MeWAY	
5-HT1B	[11C]CLMI (agonist) [11C]AZ10419369 [11C]P943	[18F]MPPF	
5-HT2A	[11C]MDL 1000907	[18F]Altanserin [18F]Altanserin-d2	
5-HT4	[11C]SB-207145		
5-HT6	[11C]GSK-215083		

Table 5 (continued)

Targets	Carbon-11 labeled	Fluorine-18 labeled	Comments
mGluR1		[18F]FTM	
mGluR5	[11C]SP 203 [11C]ABP688	[18F]SP 203 [18F]-FPFB 2-[18F]F-A-85380 (2-[18F]FA) 6-[18F]FA [18F]Nifene (agonist) [18F]AZAN [11C]CHBA-1001 [18F]ASEM [18F]SPA-RQ [18F]MK-0999 ([18F]FE-SPA-RQ) [18F]GE-179	
Nicotinic (α4β2)			
Nicotinic (α7)			
NK1			
NMDA			
NOP	[11C]NOP-1A		
Opiate (DOR)	[11C]Methylnaltrindole		
Opiate (MOR)	[11C]Diprenorphine [11C]Carfentanil (agonist) [11C]CLY2795050 (antagonist) [11C]SA4503	[18F]Fluorocetyl-diprenorphine	
Sigma 1			
Transporters			
DAT	[11C]PE21 [11C]Methylphenidate	[18F]FP-CIT [18F]FE-PE21 [18F]FECIT [18F]CFP/PB	
Glycine T1	[11C]CFpyPB [11C]GSK 9311 45		
NET	[11C]RO5013853 [11C]MeNER-d2	[18F]MeNER-d2	
SERT	[11C]DASB [11C]MADAM [11C]AFM		
TSP0	[11C]RO PK 11195 [11C]PBR28 [11C]DAA1106 [11C]DPA-713	[18F]FBR [18F]FEPPA [18F]PBI11 [18F]DPA-714	Commonly referred to as a marker of microglia activation, but target sensitivity to changes in cell number versus activation state remain unclear.
VMAT2	[11C]ERT76 [11C]DTGZ [11C]MTBZ	[18F]lorth benzamine [18F]AV-133 [18F]F-DTBZ	
Other			
	[15O]Oxygen [15O]water [11C]leucine [11C]UCB-J	[18F]FDG	Glucose utilization Oxygen utilization Blood flow Protein synthesis Marker of synaptic density
SV2a		[18F]BCPP-FE	Mitochondrial complex I density
MCI			

Table 5 Representative examples of radiotracers for CNS applications that have shown utility in humans

Targets	Carbon-11 labelled	Fluorine-18 labeled	Comments
Misfolded proteins			
β -Amyloid	[11C]PIB	[18F]Flutemetamol [18F]Florbetapir([18F]AV-45) [18F]AZD 4694 [18F]FBM [18F]FDDNP [18F]-SMIBR-W372 ([F-18]-W372) [18F]Florbetaban [18F]MK3328	
Tau		[18F] T807 (AV1451; Flortaucipir) [18F]GTP1 [18F]RO6958948 [18F]MK6240 [18F]PI-2620	Relative sensitivity to 3-repeat to 4-repeat tau isoforms remains to be confirmed.
TSPO	[11C](R)-PK 11195 [11C]PBR28 [11C]DAA1106 [11C]DPA-713 [11C]ER176	[18F]FBR [18F]FEPPA [18F]PBR111 [18F]DPA-714	Commonly referred to as a marker of microglia activation, but target sensitivity to changes in cell number versus activation state remain unclear.
SV2a	[11C]UCB-J	[18F]FDG	Marker of synaptic density Glucose utilization
	[15O]water		Blood flow

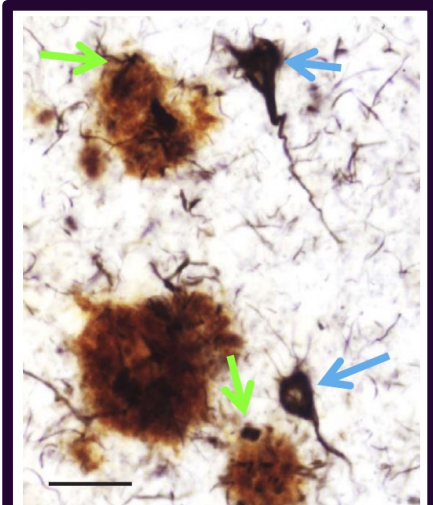
Table 5 Representative examples of radiotracers for CNS applications that have shown utility in humans

Targets			
Misfolded proteins			
β -Amyloid			
Tau			4-repeat formed.
TSPO			r of sensitivity
	[11C]DAAT106 [11C]DPA-713 [11C]ER176	[18F]PBR111 [18F]DPA-714	to changes in cell number versus activation state remain unclear.
SV2a	[11C]UCB-J	[18F]FDG	Marker of synaptic density Glucose utilization
	[15O]water		Blood flow

See past Neuroimaging PIA On-Demand Webinars on Amyloid, Tau, FDG, and TSPO PET Imaging

Alzheimer's Disease Pathology

- A = Amyloid-beta plaques
- T = Tau neurofibrillary tangles

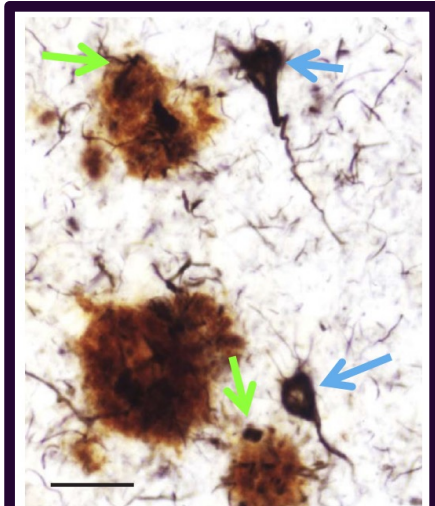


Immunohistochemistry:
A β (brown) and Tau (black)

Alzheimer's Disease Pathology

- A = Amyloid-beta plaques
- T = Tau neurofibrillary tangles

PiB binds to insoluble, fibrillar beta-amyloid aggregates

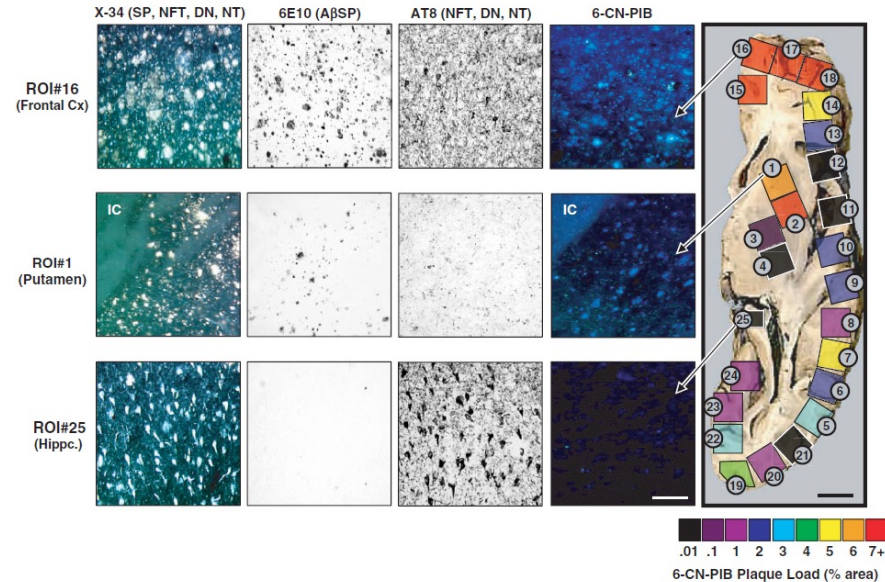


Immunohistochemistry:
Aβ (brown) and Tau (black)

Table 2 6-CN-PiB (10 μM) labeling intensity in different plaque types and amyloid-containing structures in 4% paraformaldehyde fixed, 40 μM thick tissue sections

Plaque type	6-CN-PiB intensity	X-34 intensity
Compact/coiled (NC, PhG, Crbl)		
Neuritic	++++	++++
Non-neuritic	++++	++++
Diffuse		
Amorphous (NC, PhG)	++	++
Cloud-like (Str)	++	++
Fleecy (Crbl)	0	++
Non-plaque amyloid		
Vascular	++++	++++
Neurofibrillary tangles iNFT	+ ^a	++++
Neurofibrillary tangles eNFT	+++	++++
Neuropil threads	0	++++
Dystrophic neurites	0	++++

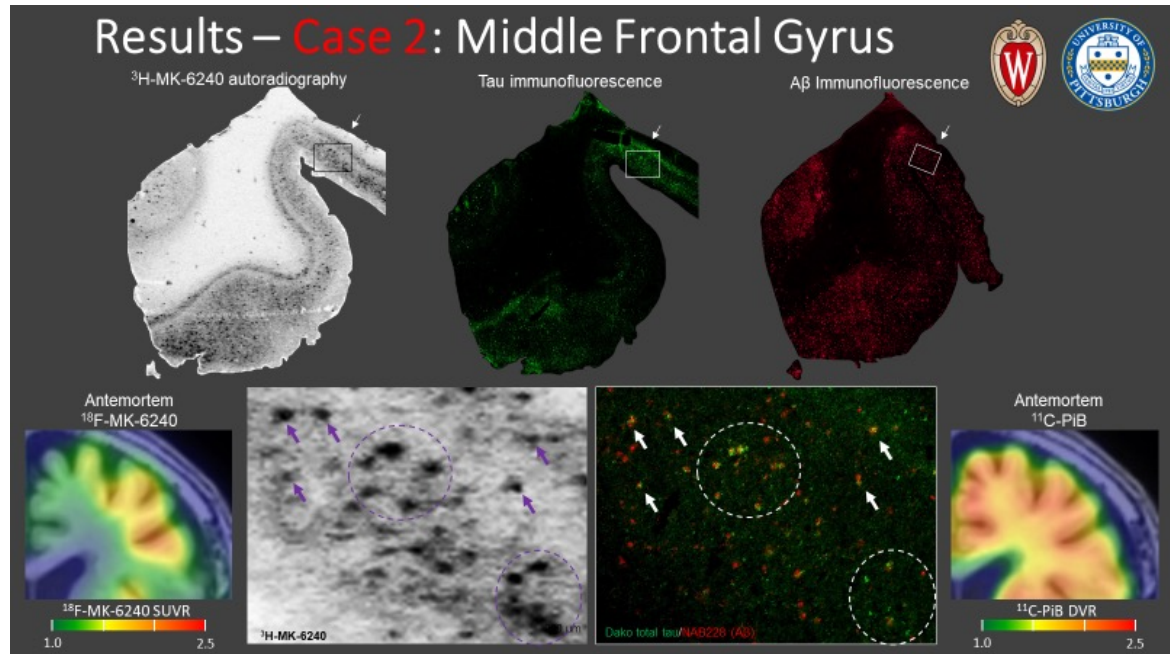
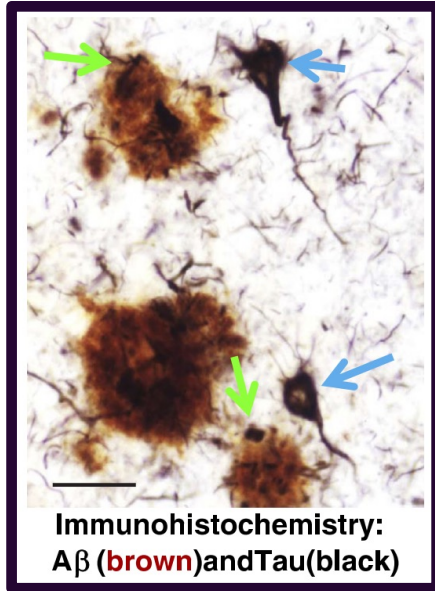
0 = no 6-CN-PiB fluorescence signal; + = very light fluorescence barely above background; ++ = light fluorescence; +++ = moderate fluorescence; ++++ = intense fluorescence.
 NC = neocortex; PhG = parahippocampal gyrus; Str = striatum; Crbl = cerebellum; iNFT = intracellular NFT; eNFT = extracellular NFT.
^aOnly a small proportion of tangles per section detected in entorhinal cortex and subiculum.



Alzheimer's Disease Pathology

- A = Amyloid-beta plaques
- T = Tau neurofibrillary tangles

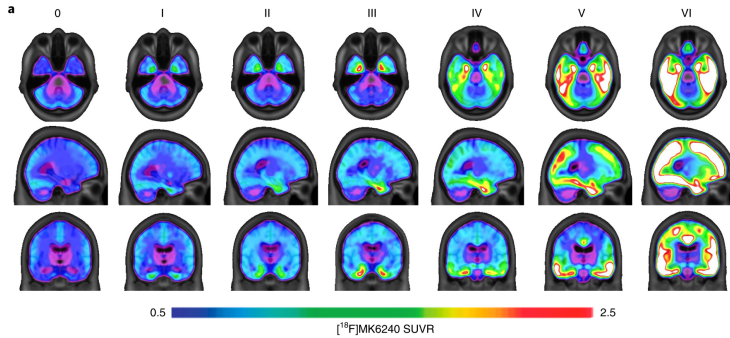
MK-6240 binds to insoluble tau aggregates: neurofibrillary tangles, neuritic pathology, neuritic plaques



Nelson, et al. *J Neuropathol Exp Neurol*. 2012

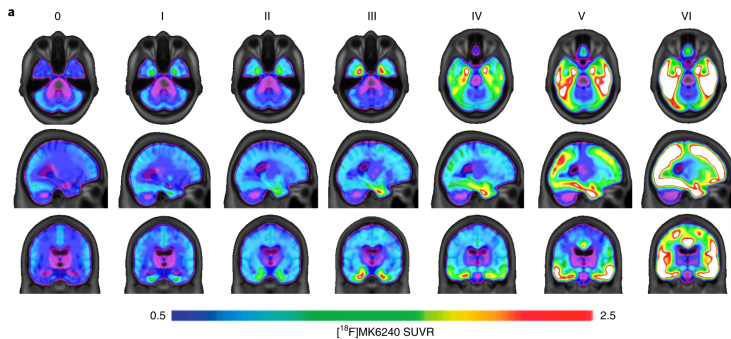
Beththausen, Ikonovic, et al. HAI 2023 (unpublished)

Tau PET imaging recapitulates Braak neuropathological staging



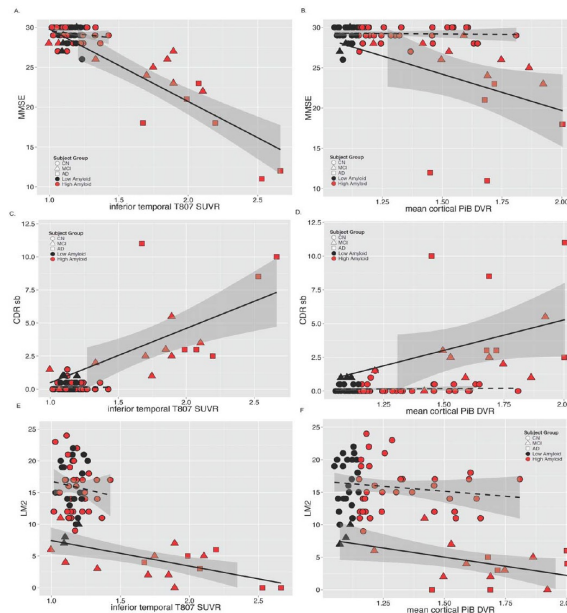
Therriault, et al. Nature Aging, 2022

Tau PET imaging recapitulates Braak neuropathological staging



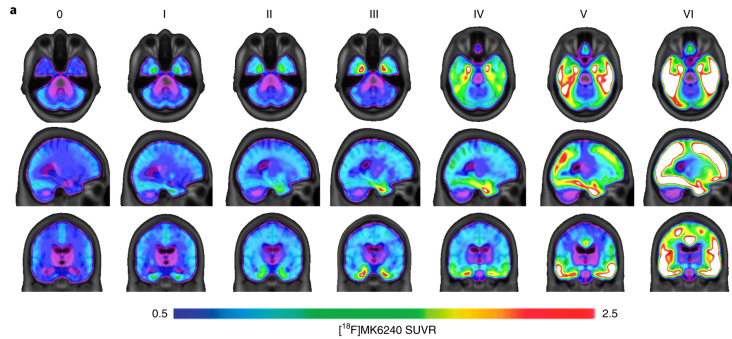
Therriault, et al. Nature Aging, 2022

PET-measured AD pathology, especially tau, associates with cognitive deficits



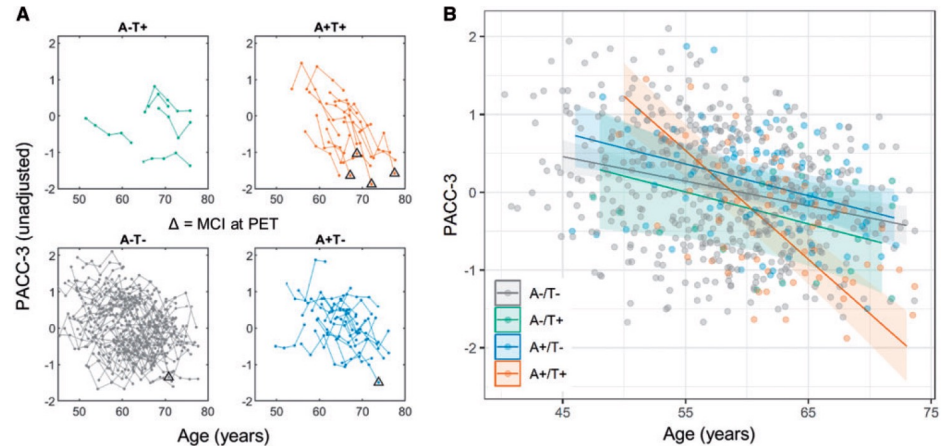
Johnson KA, et al. Annals of Neurology, 2015

Tau PET imaging recapitulates Braak neuropathological staging

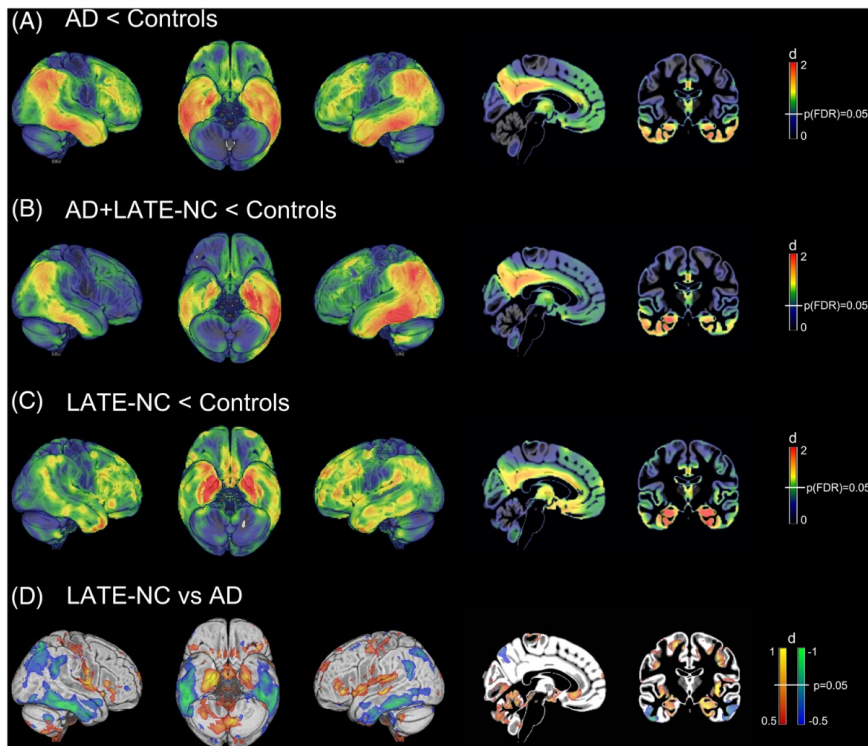


Therriault, et al. Nature Aging, 2022

A+T+ associates with accelerated preclinical cognitive decline



Betthausen, et al. Brain, 2020



FDG hypometabolic spatial patterns may indicate underlying neuropathology

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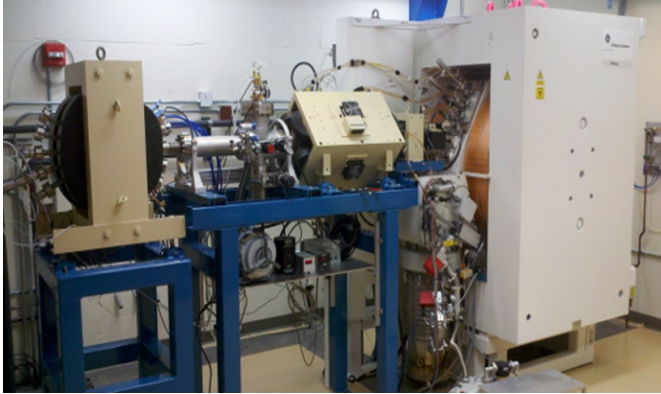
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Introduction – Basics of PET

- How is PET different from other imaging modalities?
- What is a PET tracer?
- **How do we get a PET image?**
- How do we quantify PET?

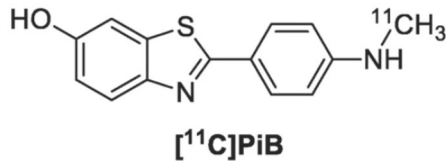
PET Scanning Overview

Cyclotron



Radiochemical
Synthesis
and Purification

PET Tracer



Administer

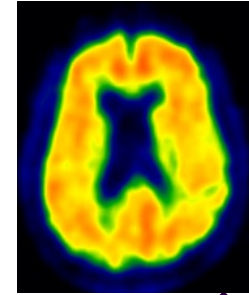
- IV Injection
- Inhalation



PET Steps:

- 1) Isotope Production
- 2) Synthesize
- 3) Administer
- 4) Scan
- 5) Process

Image Processing



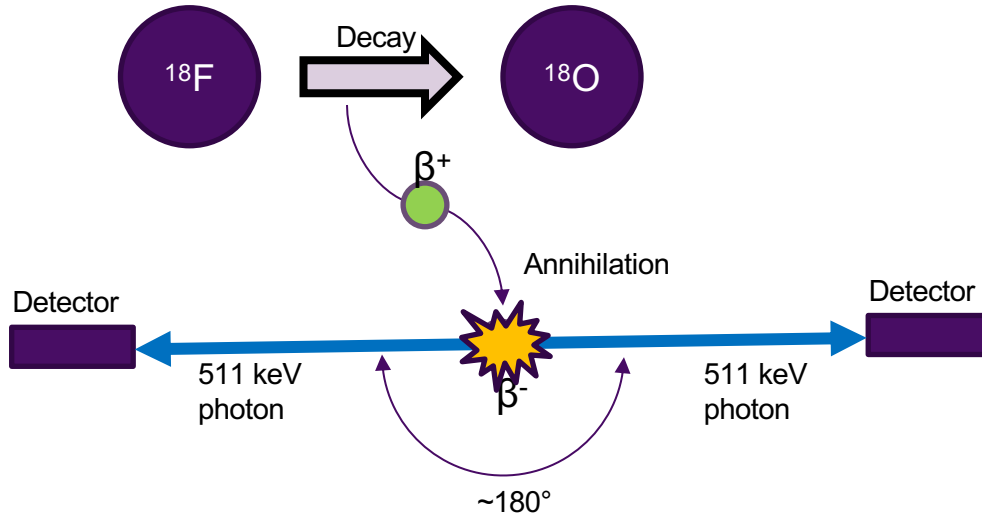
Scan



PET/CT Scanner

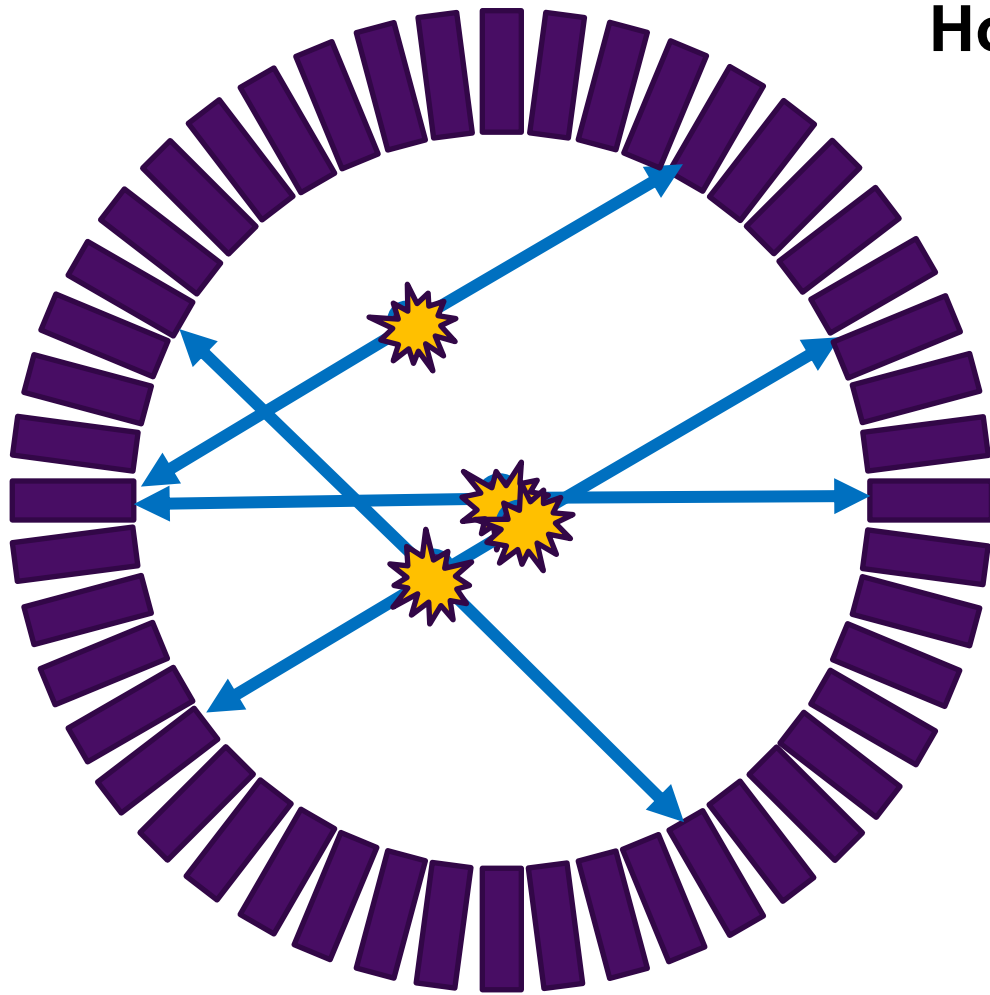
How to make a PET image

Radioactive Decay and Positron Annihilation

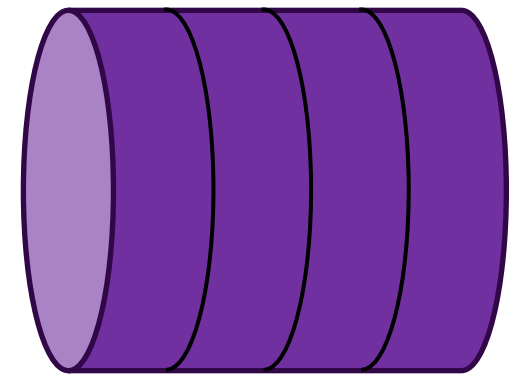


- 1) Radioisotope decays
- 2) Positron annihilation
- 3) "Coincident" Photons detected

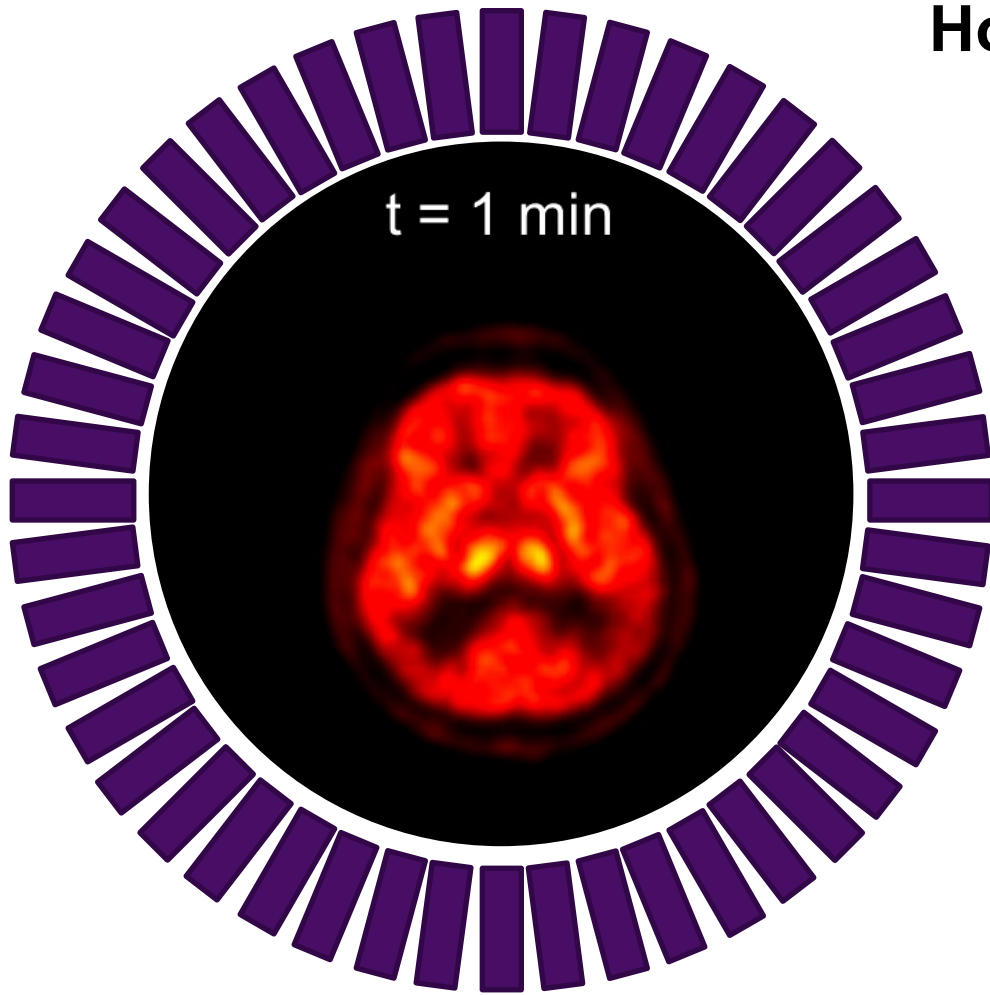
How to make a PET image



- 1) Radioisotope decays
- 2) Positron annihilation
- 3) "Coincident" Photons detected
- 4) Detect many events over time

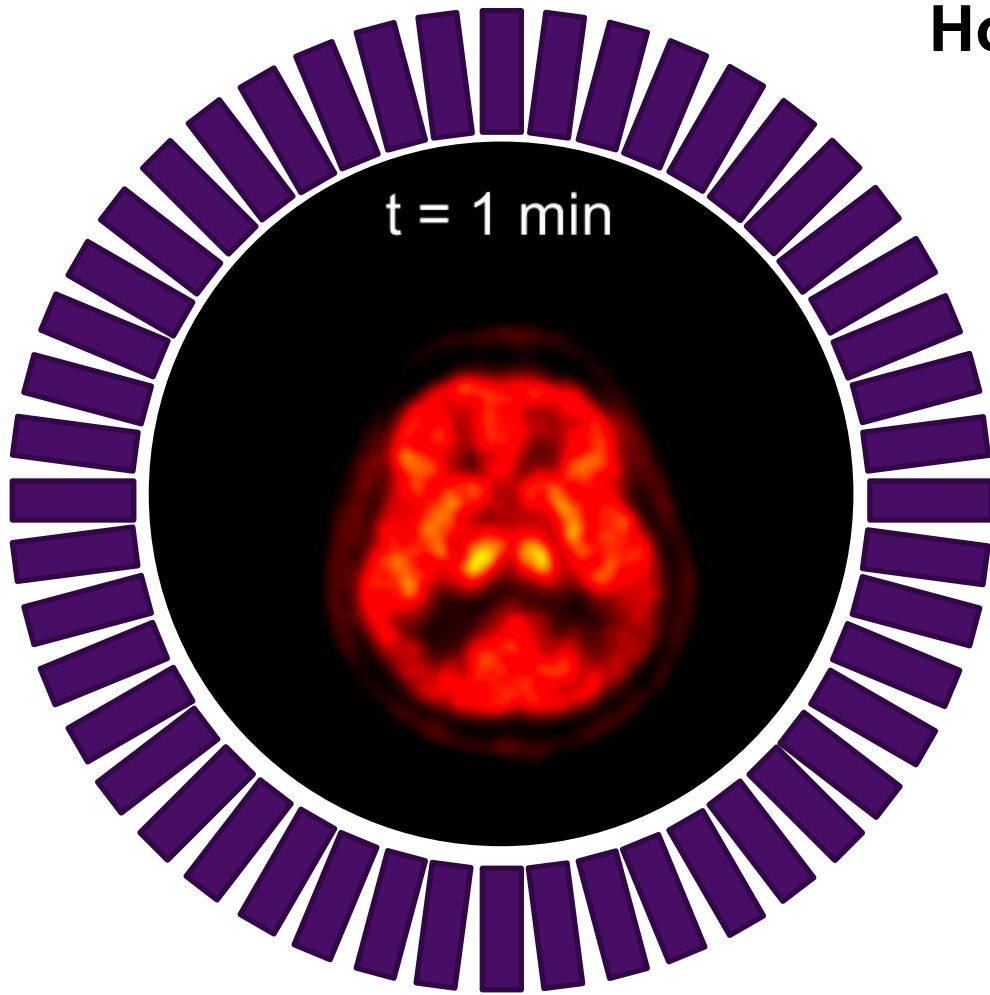


How to make a PET image

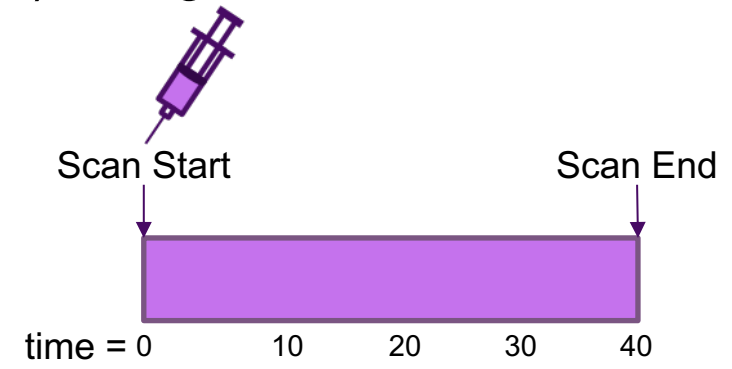


- 1) Radioisotope decays
- 2) Positron annihilation
- 3) "Coincident" Photons detected
- 4) Detect many events over time
- 5) Image Reconstruction

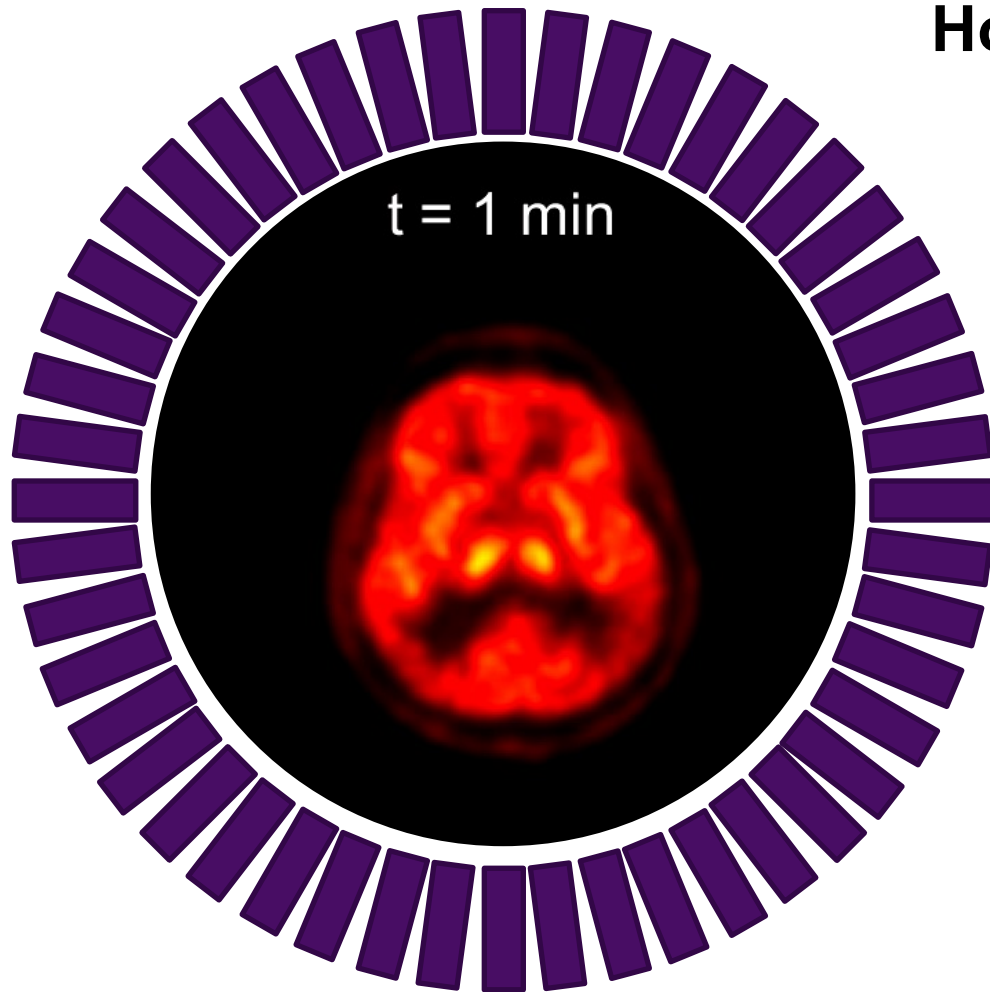
How to make a PET image



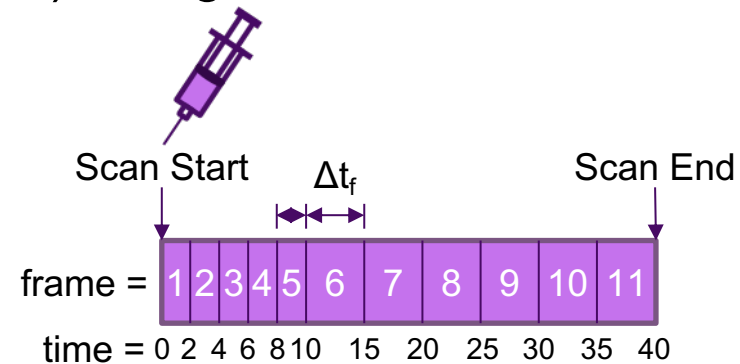
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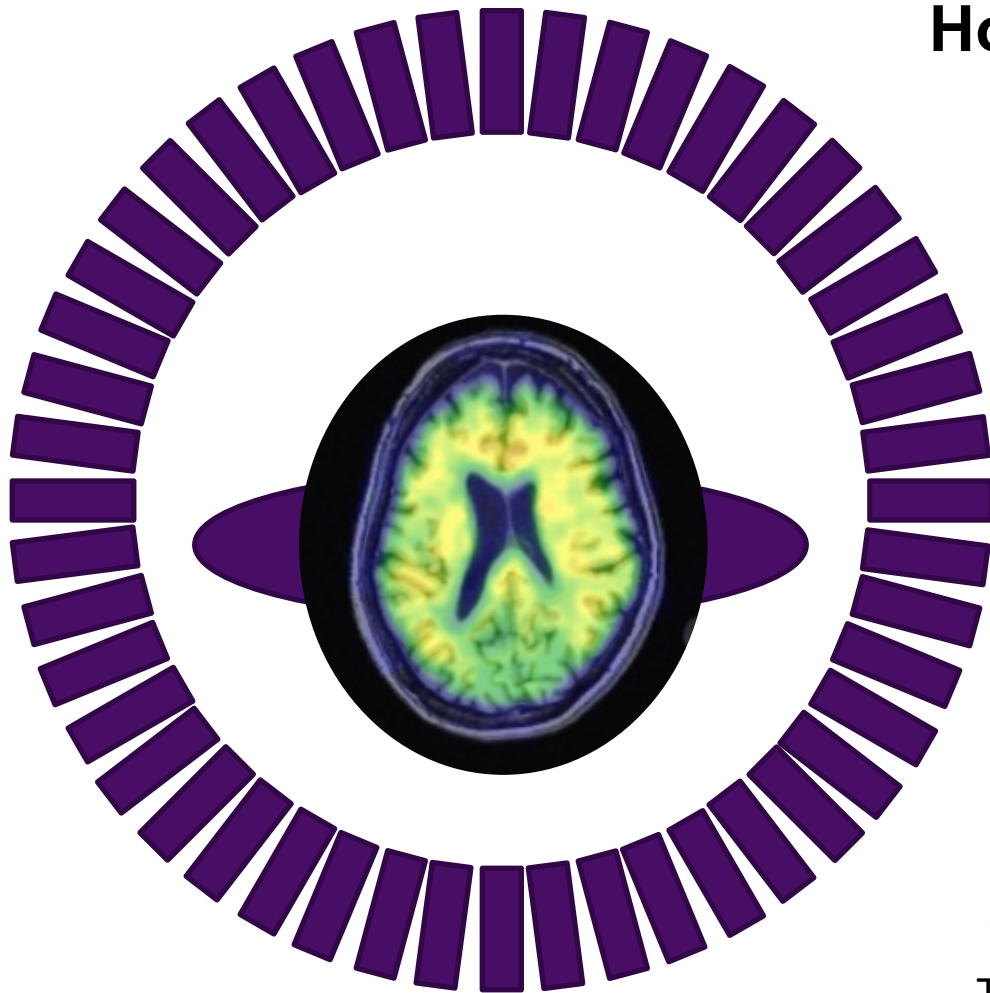
How to make a PET image



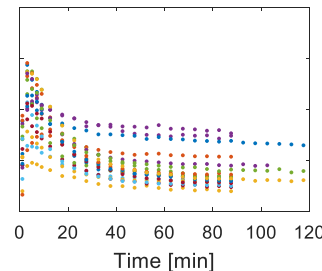
- 1) Radioisotope decays
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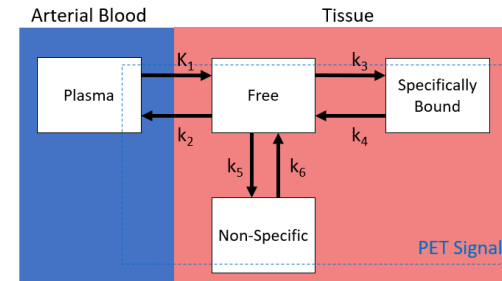
How to make a PET image



- 1) Radioisotope decays
- 2) Positron annihilation
- 3) “Coincident” Photons detected
- 4) Detect many events over time
- 5) Image Reconstruction
- 6) Image Processing



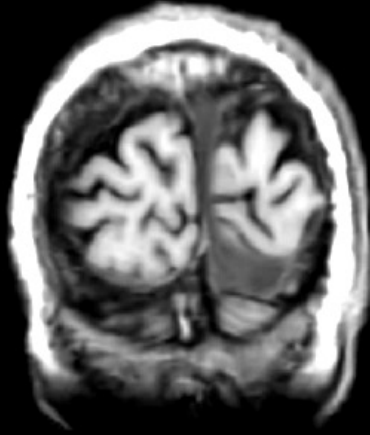
Time-Activity Curve



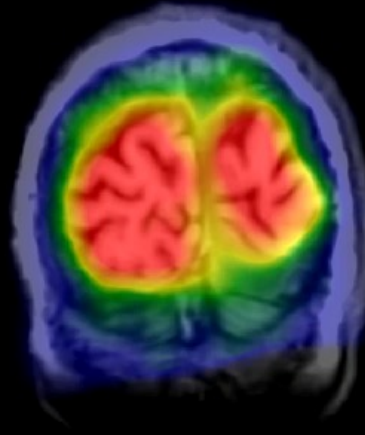
Kinetic Analysis

Example Parametric Images

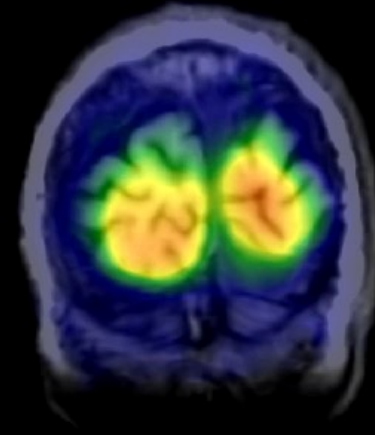
T1 MRI



MK-6240 SUVR



PiB DVR



PET provides macroscopic quantitative measures of underlying molecular biology and/or physiology *in vivo*

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Introduction – Basics of PET

- How is PET different from other imaging modalities?
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- How do we get a PET image?
- **How do we quantify PET?**

Optimal Quantification Method is a Trade-off

Experimental
Complexity



Accuracy and
Information



Optimal Quantification Method is a Trade-off

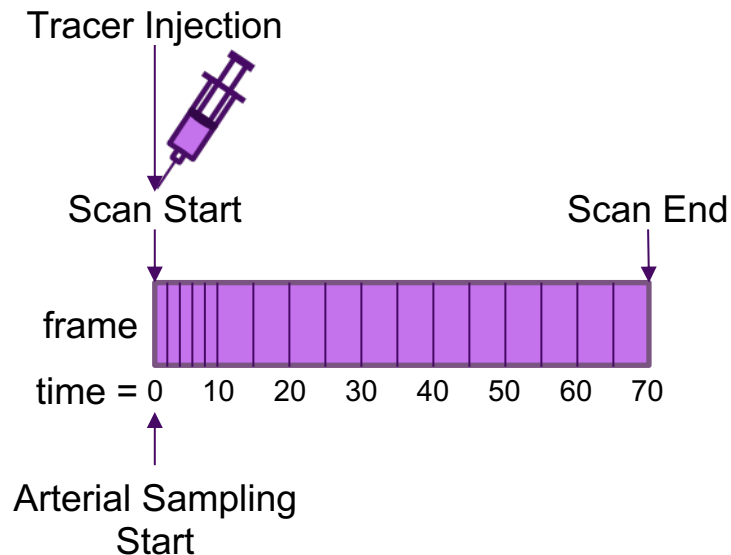
Experimental Complexity



PET with arterial sampling

- arterial cannulation
- long scan duration
- + full kinetic modeling

Accuracy and Information



Optimal Quantification Method is a Trade-off

Experimental Complexity



PET with arterial sampling

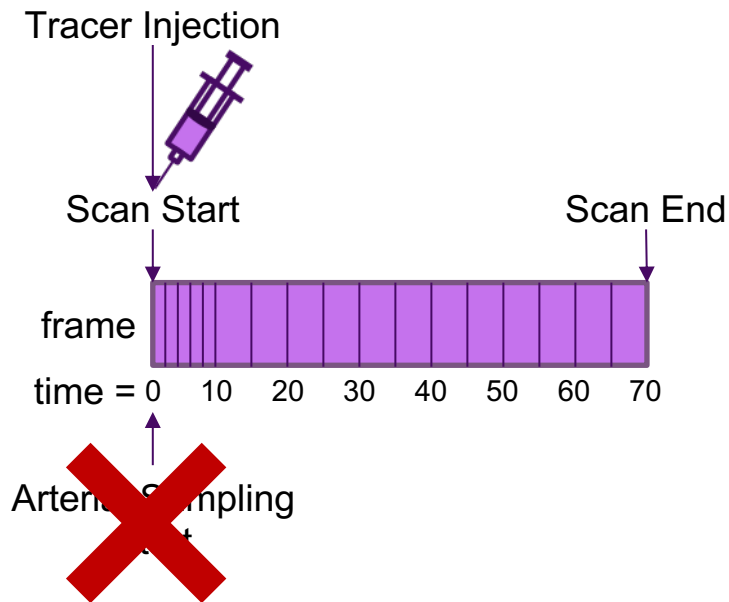
- arterial cannulation
- long scan duration
- + full kinetic modeling

Full-Dynamic Imaging

- + no arterial sampling
- long scan duration
- + quantitative accuracy



Accuracy and Information



Optimal Quantification Method is a Trade-off

Experimental Complexity



PET with arterial sampling

- arterial cannulation
- long scan duration
- + full kinetic modeling

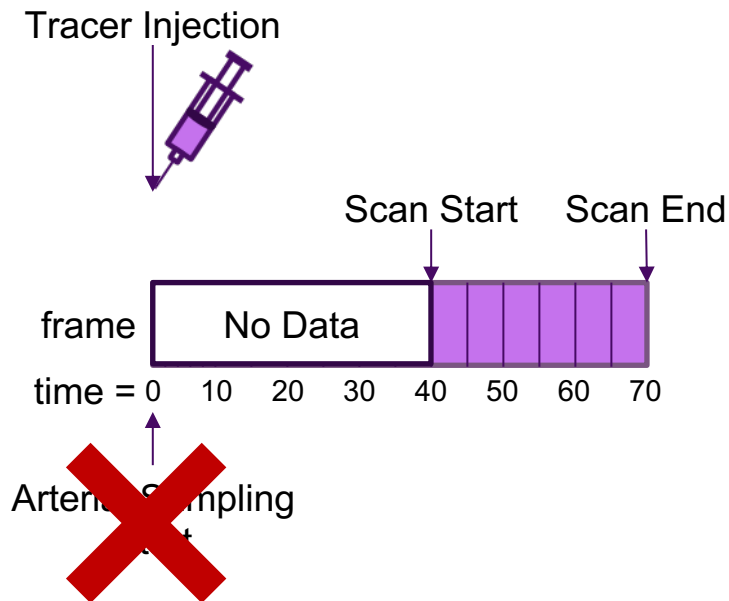
Full-Dynamic Imaging

- + no arterial sampling
- long scan duration
- + quantitative accuracy

Late-frame Dynamic Imaging

- less accurate
- binding estimates impacted by blood flow
- + short scan duration
- + some kinetic information

Accuracy and Information



Experimental
Complexity**PET with arterial sampling**

- arterial cannulation
- long scan duration
- + full kinetic modeling

Full-Dynamic Imaging

- + no arterial sampling
- long scan duration
- + quantitative accuracy

Late-frame Dynamic Imaging

- less accurate
- binding estimates impacted by blood flow
- + short scan duration
- + some kinetic information

Late-frame Static Imaging

- less accurate
- no kinetic information
- + short scan duration
- + less data (smaller files)

Accuracy and
Information

Tracer Injection



Scan Start

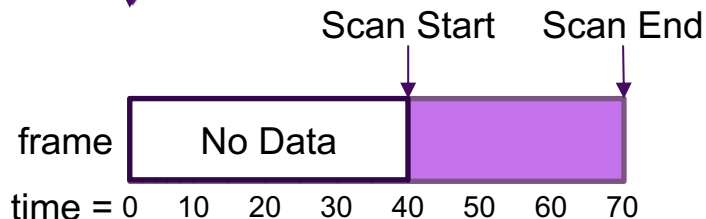
Scan End

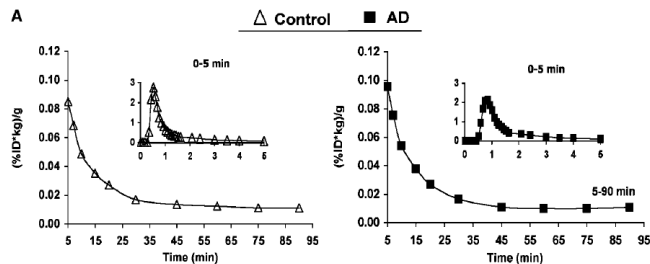
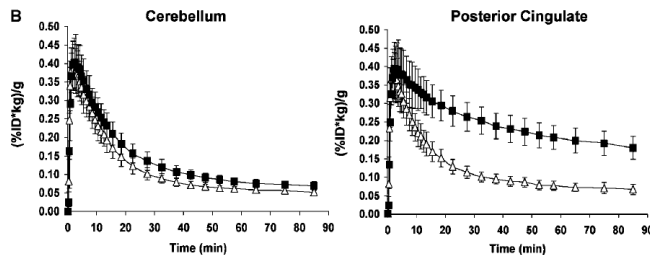
frame

No Data

time = 0 10 20 30 40 50 60 70

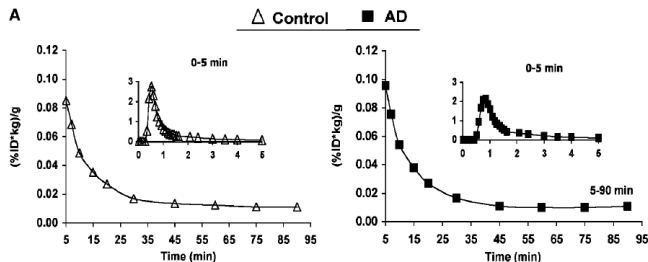
Arterial Sampling



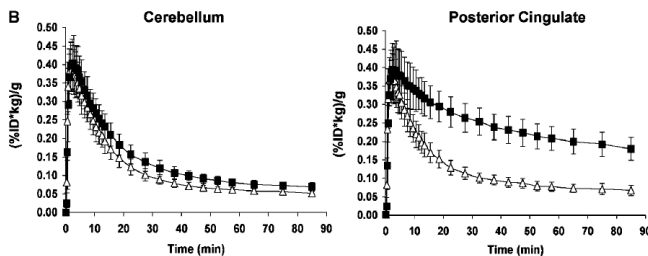
Arterial
Input FunctionBrain
Time-Activity
Curves

Adapted from Price, et al. JCBFM. 2005

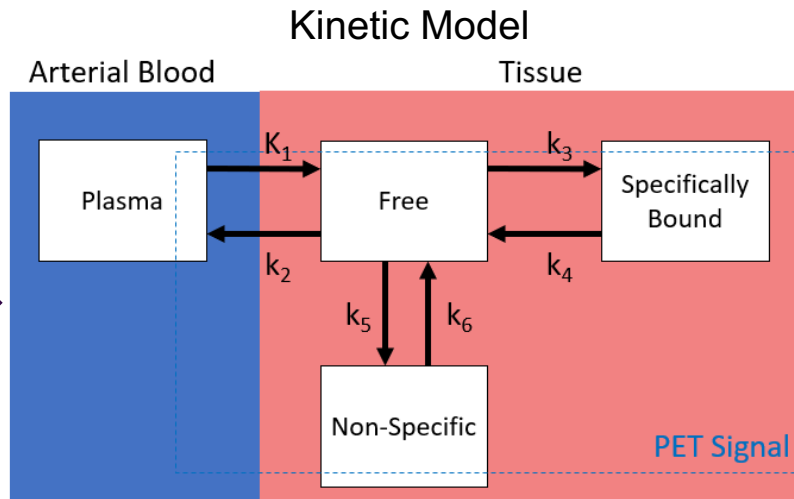
Arterial
Input Function



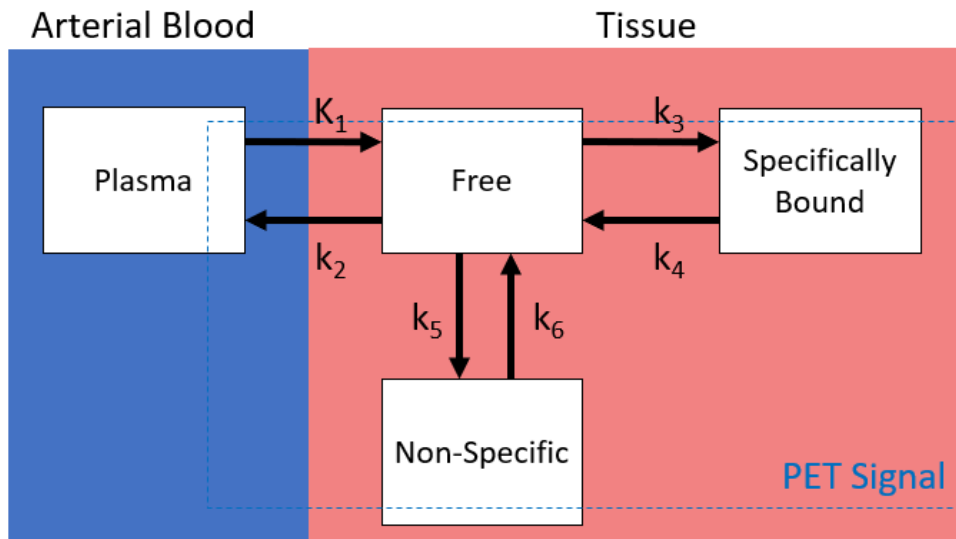
Brain
Time-Activity
Curves



Adapted from Price, et al. JCBFM. 2005



Kinetic Model

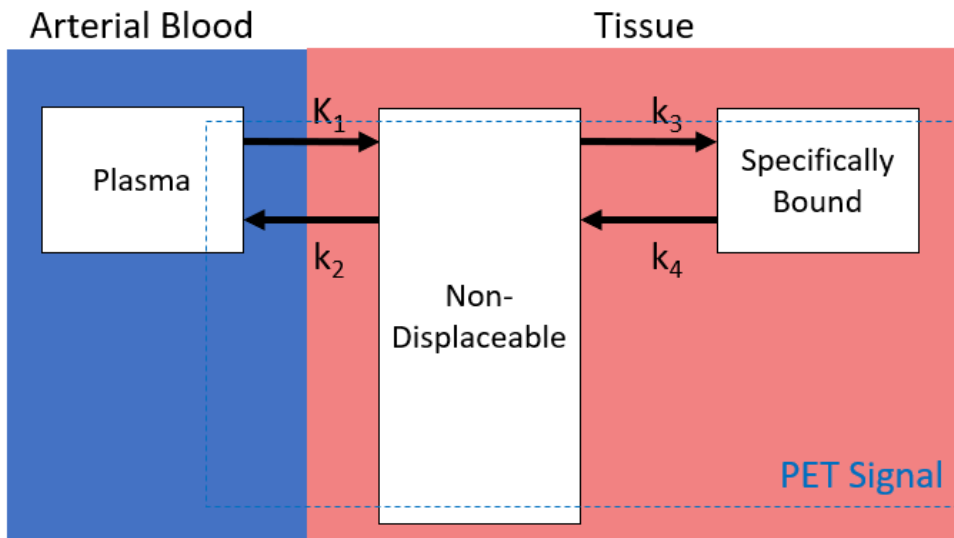


Distribution Volume

$$V_T = \frac{C_T}{C_P}$$

$$V_T = V_F + V_{NS} + V_S$$

2-tissue Compartment Model

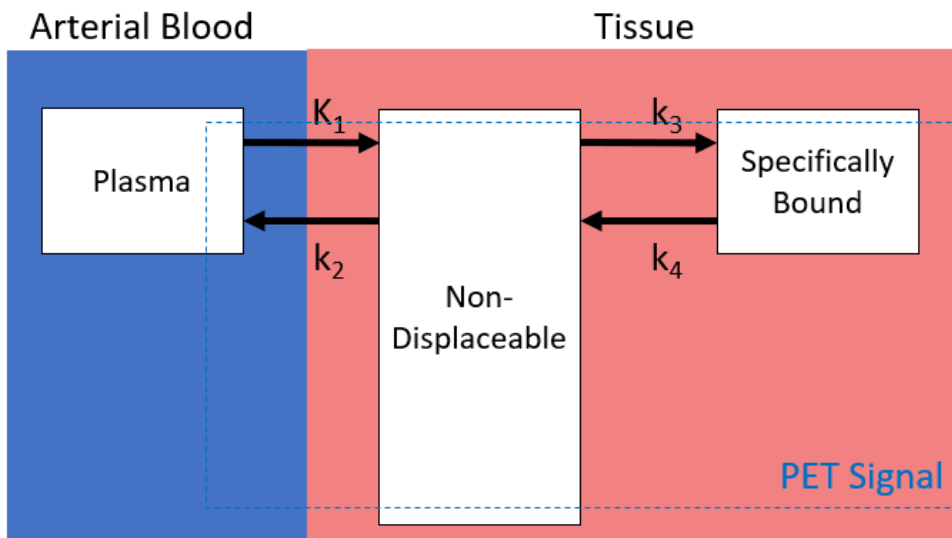


Distribution Volume

$$V_T = \frac{C_T}{C_P}$$

$$V_T = V_{ND} + V_S$$

2-tissue Compartment Model



B_{max} is the density of the target receptor

$1/K_D$ is referred to as the affinity

Distribution Volume

$$V_T = \frac{C_T}{C_P}$$

$$V_T = V_{ND} + V_S$$

Distribution Volume Ratio (DVR)

$$\frac{V_T^{target}}{V_T^{ref}} = \frac{V_{ND} + V_S}{V_{ND}}$$

$$DVR = 1 + BP_{ND}$$

$$BP = \frac{B_{max}}{K_D}$$

2-tissue Compartment Model

Arterial Blood

Tissue

Distribution Volume

 k_1 k_2

$$V_T = \frac{C_T}{C_A}$$

For reversibly bound PET ligands, **Binding Potential** (and therefore DVR) is a quantitative in vivo measure that is directly proportional to molecular receptor density

B_{max} is the density of the target receptor

$1/K_D$ is referred to as the affinity

$$DVR = 1 + BP_{ND}$$

$$BP = \frac{B_{max}}{K_D}$$

Experimental Complexity

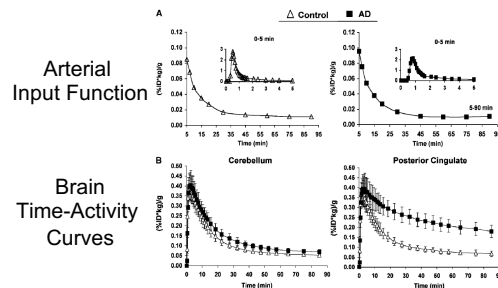


PET with arterial sampling

- V_T
- Rate Constants
- Tracer Metabolism



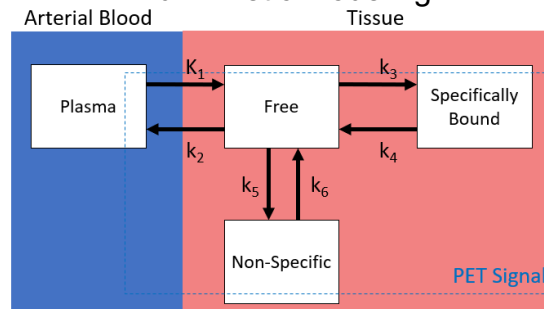
Accuracy and Information



Adapted from Price, et al. JCBFM. 2005



Full Kinetic Modeling



Optimal Quantification Method is a Trade-off

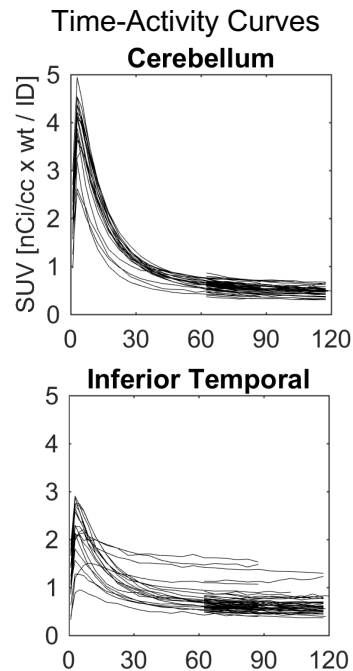
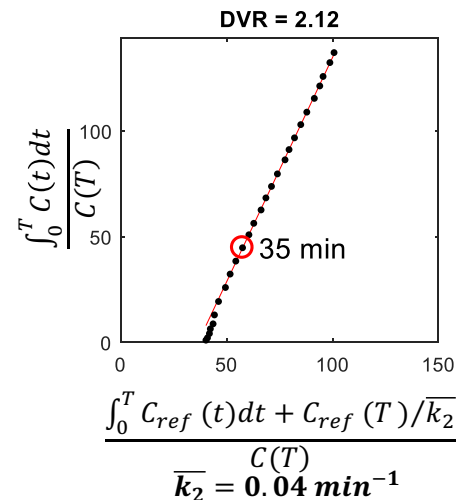
Experimental
Complexity

PET with arterial sampling

- V_T
- Rate Constants
- Tracer Metabolism

Full-Dynamic Imaging

- DVR (reference tissue methods)
- R_1 (relative perfusion)

Accuracy and
InformationLogan Graphical Analysis in
AD Middle Temporal Gyrus

Experimental
Complexity

PET with arterial sampling

- V_T
- Rate Constants
- Tracer Metabolism

Full-Dynamic Imaging

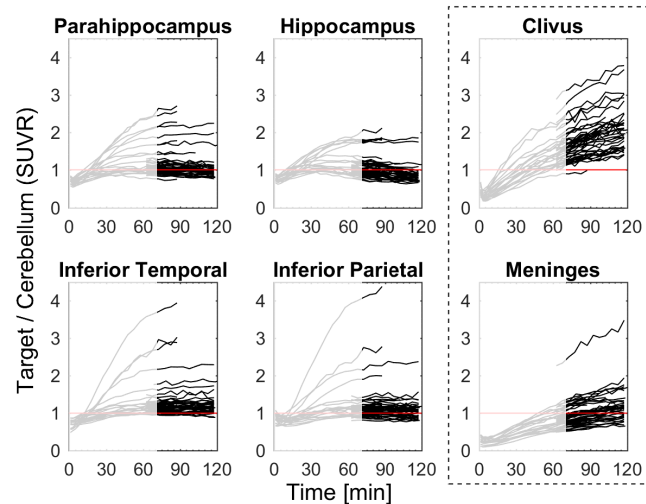
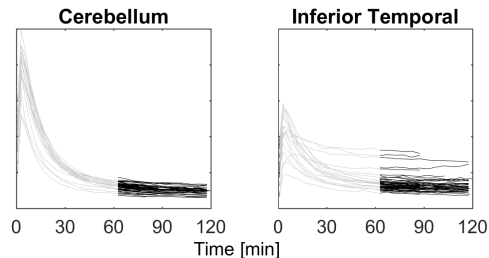
- DVR (reference tissue methods)
- R_1 (relative perfusion)

Late-frame Dynamic Imaging

- SUVR

Late-frame Static Imaging

- SUVR

Accuracy and
Information

ALZHEIMER'S  ASSOCIATION®

AAIC>23

PET Image Processing

- MR-Guided Image Processing
- PET only Image Processing
- Other Considerations

ALZHEIMER'S  ASSOCIATION®

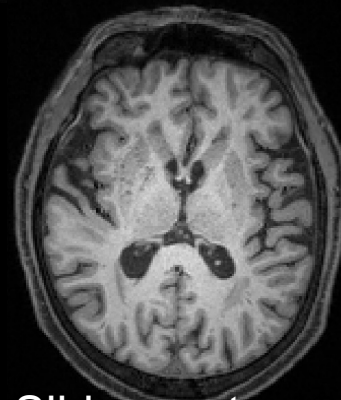
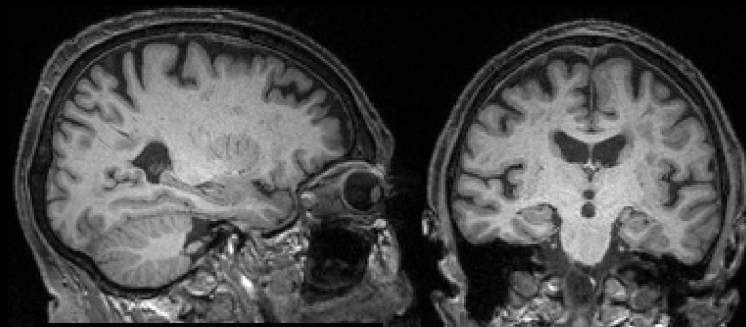
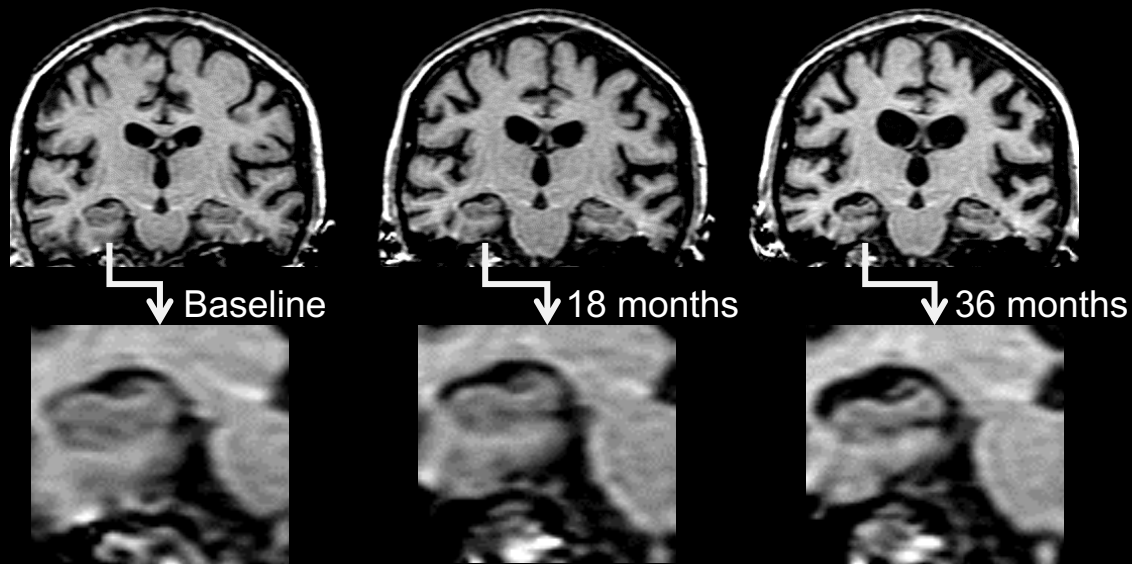
AAIC>23

PET Image Processing

- **MR-Guided Image Processing**
- PET only Image Processing
- Other Considerations

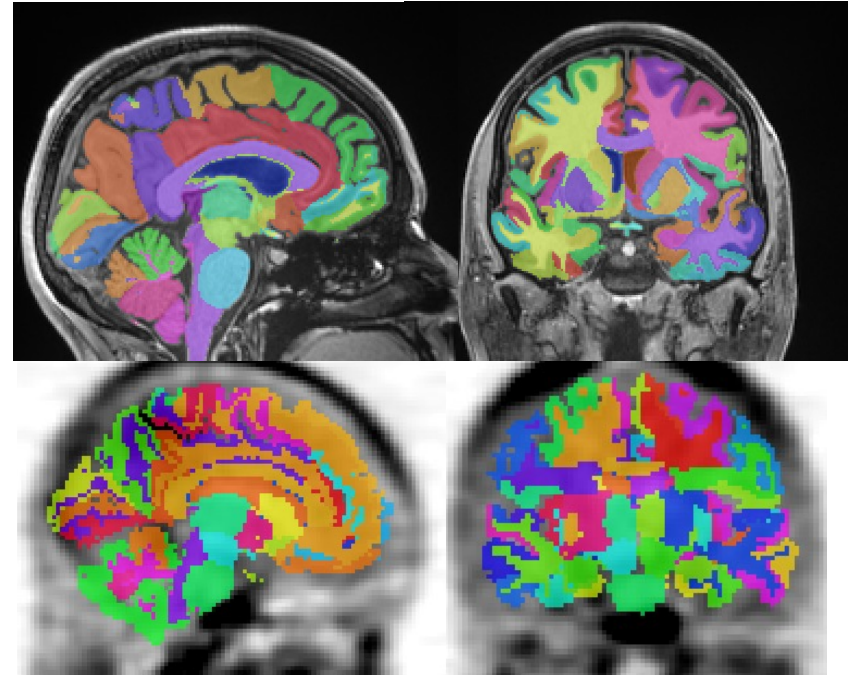
T1-weighted MRI

- High-resolution (~1 mm) information about **neuroanatomy** and **neurodegeneration**
- Can be acquired in any orientation in ~4-6 minutes
- Good contrast between different tissues (GM, WM, CSF)



Slide courtesy of Dr. Dave Cash

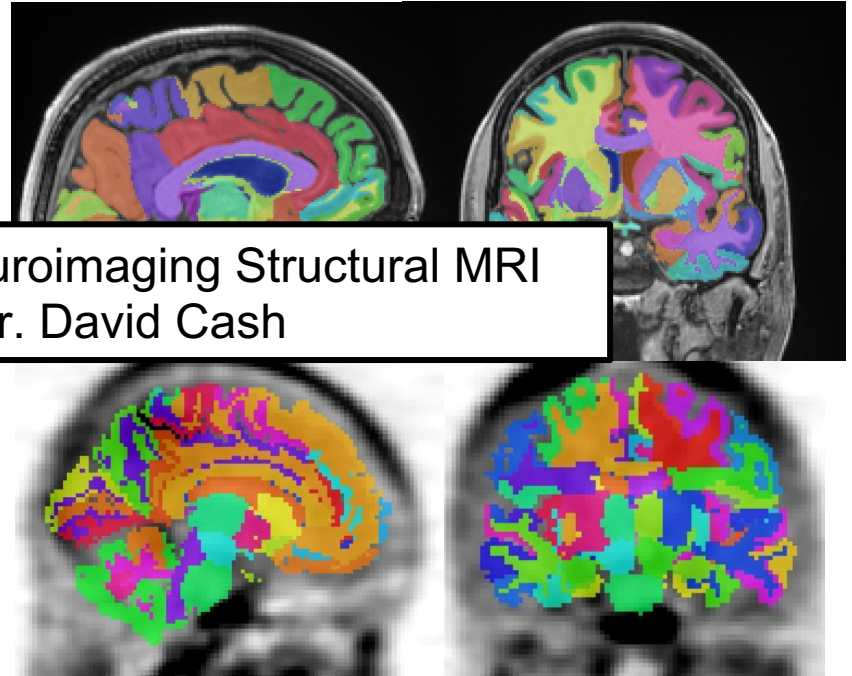
- Structural T1 provides high-resolution anatomical context for other lower resolution modalities (fMRI, DWI, PET)
- Regions of interest (ROIs) defined on the structural T1 scan can be transferred to co-registered images
- Tissue properties from segmentation can also provide some information on partial volume effect (mixture of different tissues)



Slide courtesy of Dr. Dave Cash

- Structural T1 provides high-resolution anatomical context for other lower resolution modalities (fMRI, DWI, PET)
- Regions of interest (ROIs) defined on the structural T1 co-registered
- Tissue properties from segmentation can also provide some information on partial volume effect (mixture of different tissues)

See Previous Basics of Neuroimaging Structural MRI presented by Dr. David Cash



Process MRI
(ROI Parcellation)

Smooth/De-noise

Interframe Alignment
(i.e., motion correction)

Co-Registration to MRI

Extract Reference
Region TAC

Generate Parametric
Image(s)

Process MRI
(ROI Parcellation)

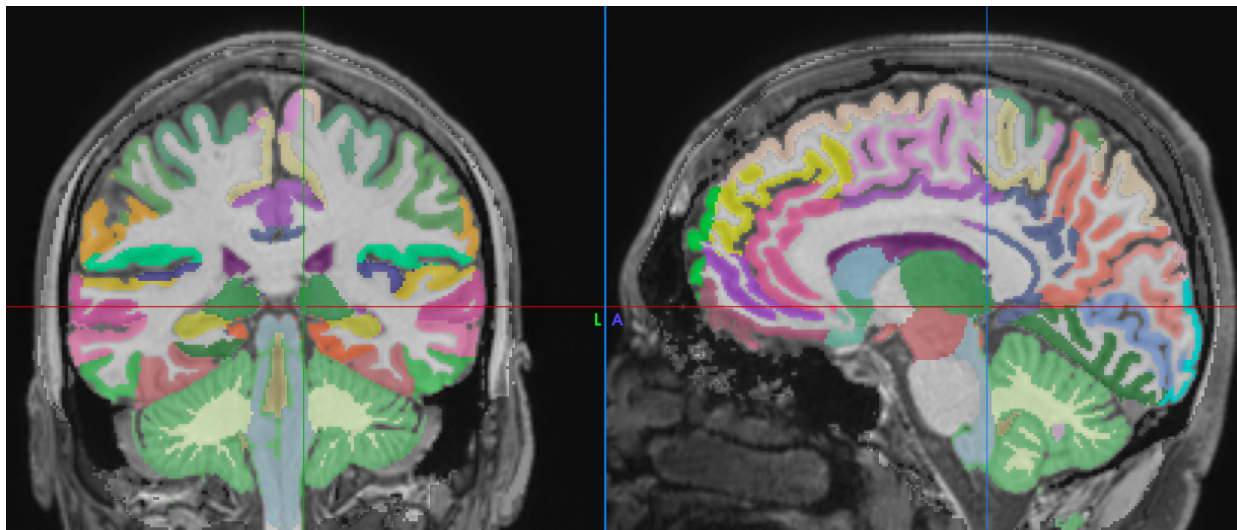
Smooth/De-noise

Interframe Alignment
(i.e., motion correction)

Co-Registration to MRI

Extract Reference
Region TAC

Generate Parametric
Image(s)



See Previous Basics of Structural MRI Webinar
presented by Dr. Dave Cash

Process MRI
(ROI Parcellation)

Smooth/De-noise

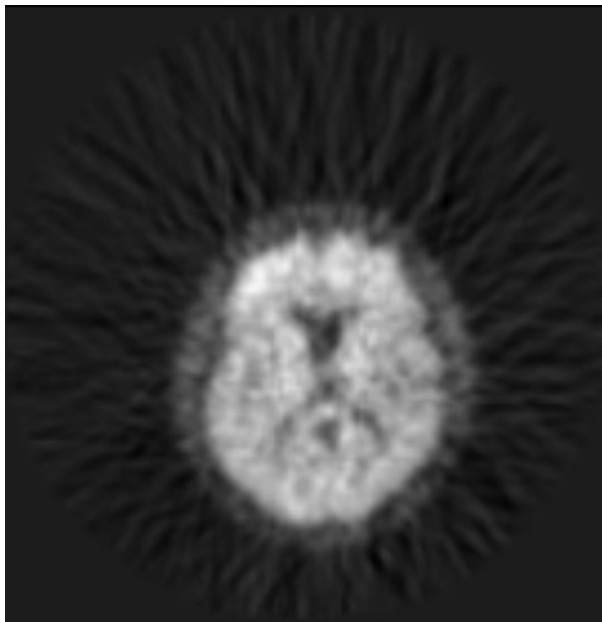
Interframe Alignment
(i.e., motion correction)

Co-Registration to MRI

Extract Reference
Region TAC

Generate Parametric
Image(s)

Individual PET frames are noisy!



Single [^{11}C]PiB PET Frame (30-35 min)

Process MRI
(ROI Parcellation)

Smooth/De-noise

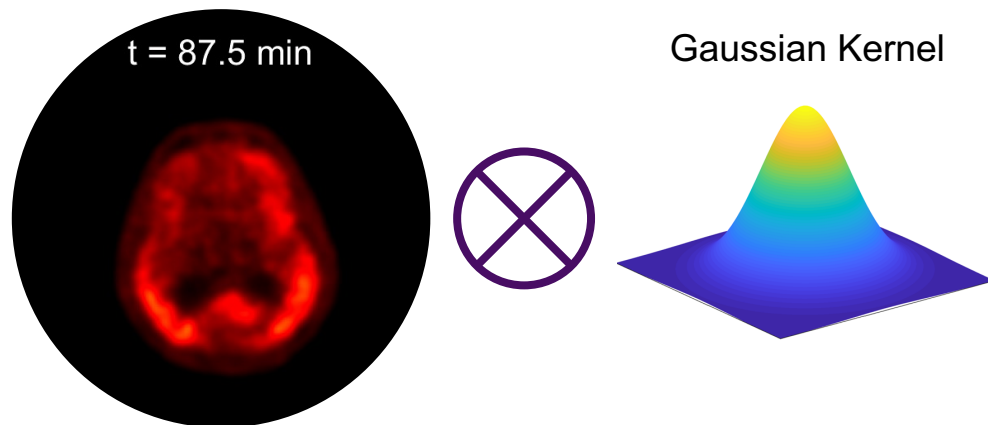
Interframe Alignment
(i.e., motion correction)

Co-Registration to MRI

Extract Reference
Region TAC

Generate Parametric
Image(s)

Spatial Smoothing



- Spatial smoothing reduces voxel variance but increase covariance of adjacent voxels
- Can be applied during or after image reconstruction
- Over smoothing reduces spatial resolution

Process MRI
(ROI Parcellation)

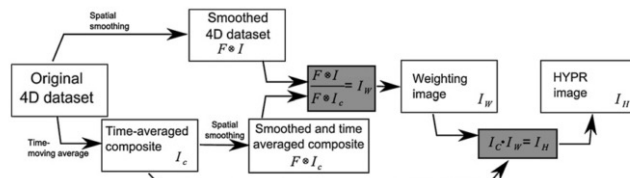
Smooth/De-noise

Interframe Alignment
(i.e., motion correction)

Co-Registration to MRI

Extract Reference
Region TAC

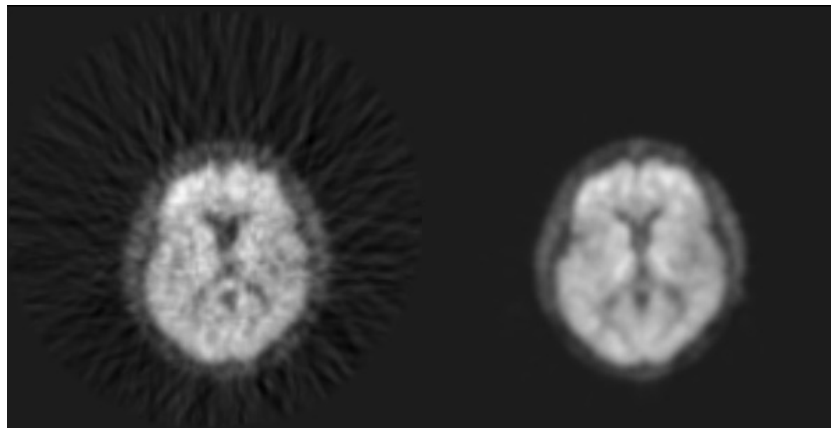
Generate Parametric
Image(s)



Christian, et al. *J Nucl Med.* 2010

3mm Smoothing

3mm Smoothing + HYPR-LR



Single [^{11}C]PIB PET Frame (30-35 min)

Process MRI
(ROI Parcellation)

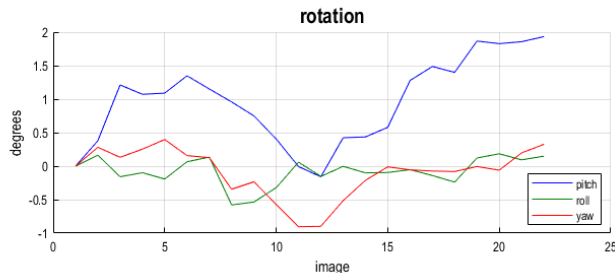
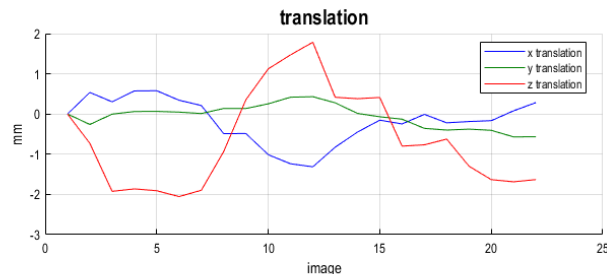
Smooth/De-noise

Interframe Alignment
(i.e., motion correction)

Co-Registration to MRI

Extract Reference
Region TAC

Generate Parametric
Image(s)



Degrees of Freedom

- Translation (x3)
- Rotation (x3)
- Shear (x3)
- Scaling/Zoom (x3)

} Rigid Body

- Correction for motion between PET frames
- Will not correct for misaligned attenuation maps
- Will not correct for motion within a frame

Process MRI
(ROI Parcellation)

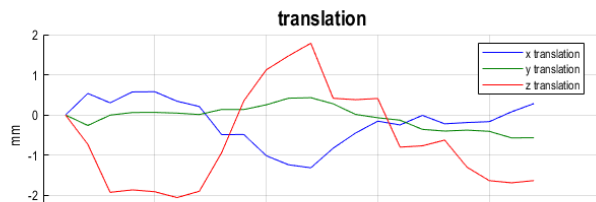
Smooth/De-noise

Interframe Alignment
(i.e., motion correction)

Co-Registration to MRI

Extract Reference
Region TAC

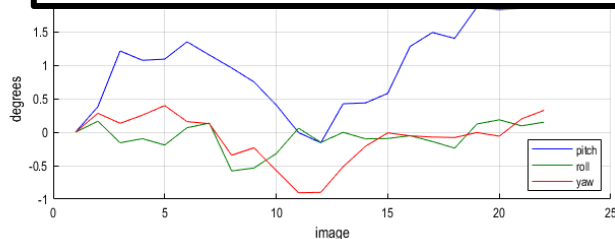
Generate Parametric
Image(s)



Degrees of Freedom

- Translation (x3)
 - Rotation (x3)
 - Shear (x3)
- } Rigid Body

See Previous Basics of Neuroimaging Data Structure and Formats presented by Dr. Ludovca Griffanti



- Correction for motion between PET frames
- Will not correct for misaligned attenuation maps
- Will not correct for motion within a frame

Process MRI
(ROI Parcellation)

Smooth/De-noise

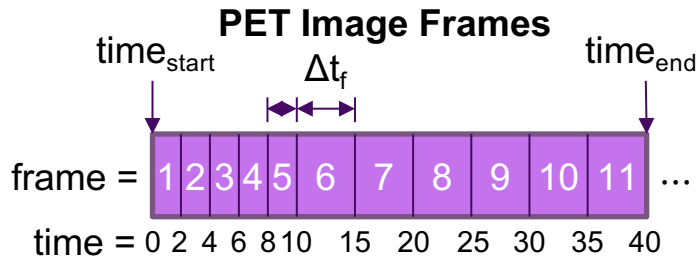
Interframe Alignment
(i.e., motion correction)

Co-Registration to MRI

Extract Reference
Region TAC

Generate Parametric
Image(s)

PET frames can be different durations



SUM PET Image

$$\frac{\sum_{f=time_{start}}^{time_{end}} C(t)_f \times \Delta t_f}{\sum_{f=time_{start}}^{time_{end}} \Delta t_f}$$

List Mode

$$\sum_{i=t_{start}}^{t_{end}} \frac{counts_i}{t_i}$$

- Start with SUM PET image (time duration-weighted average)
- Register each frame to SUM image (can do this iteratively)

*Note that an average of PET frames is equivalent to a SUM image only if all the frames are the same duration

MR-Guided PET Image Processing

Process MRI
(ROI Parcellation)

Smooth/De-noise

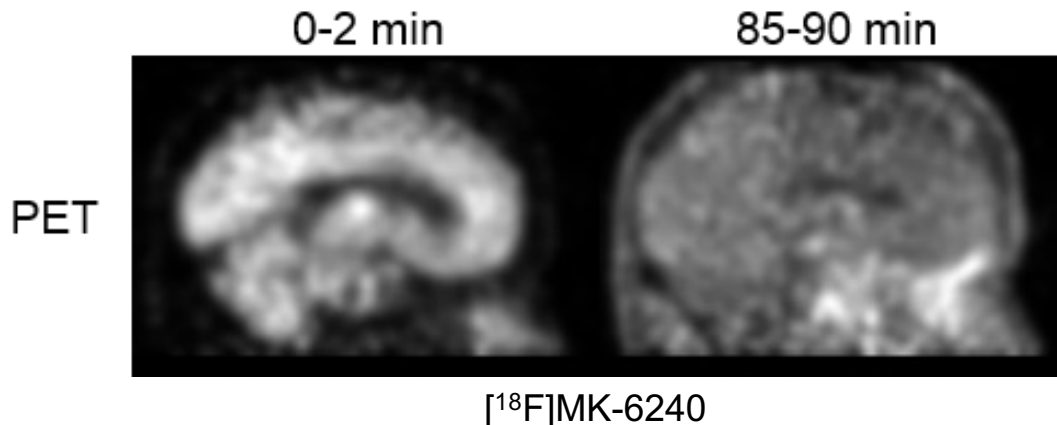
Interframe Alignment
(i.e., motion correction)

Co-Registration to MRI

Extract Reference
Region TAC

Generate Parametric
Image(s)

Spatial information can change dramatically during the scan



- Can create challenges for inter-frame alignment

MR-Guided PET Image Processing

Process MRI
(ROI Parcellation)

Smooth/De-noise

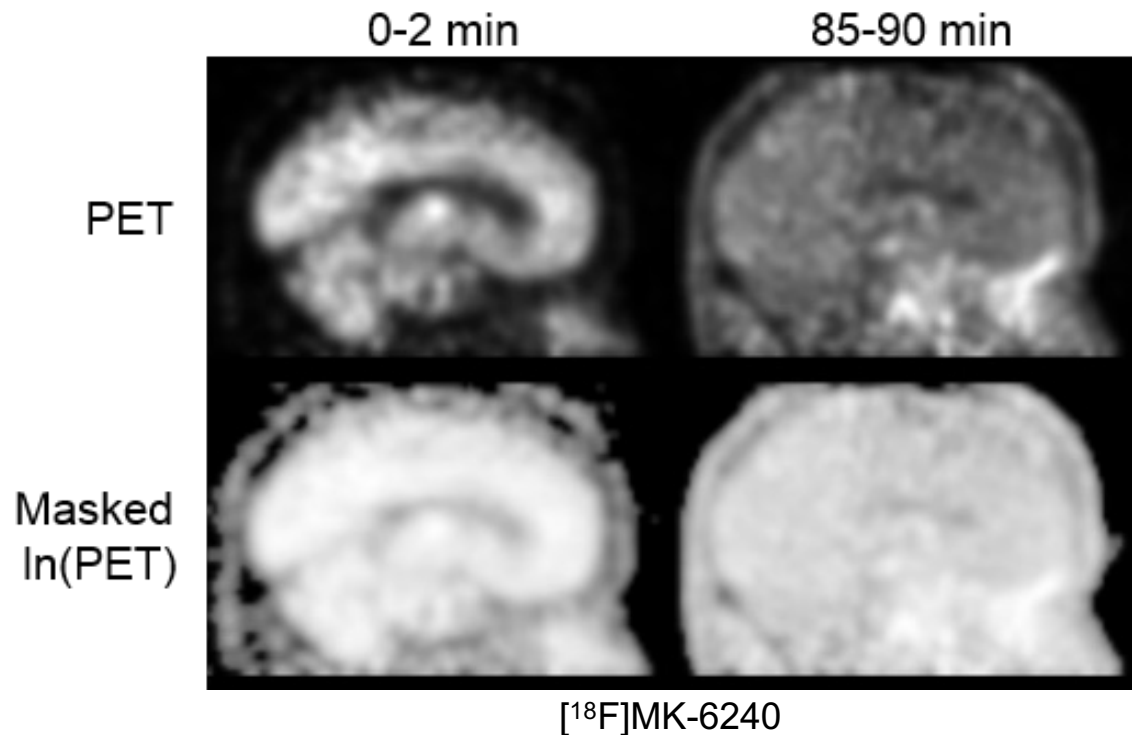
Interframe Alignment
(i.e., motion correction)

Co-Registration to MRI

Extract Reference
Region TAC

Generate Parametric
Image(s)

Image transforms can help reduce dissimilarities between frames



MR-Guided PET Image Processing

Process MRI
(ROI Parcellation)

Smooth/De-noise

Interframe Alignment
(i.e., motion correction)

Co-Registration to MRI

Extract Reference
Region TAC

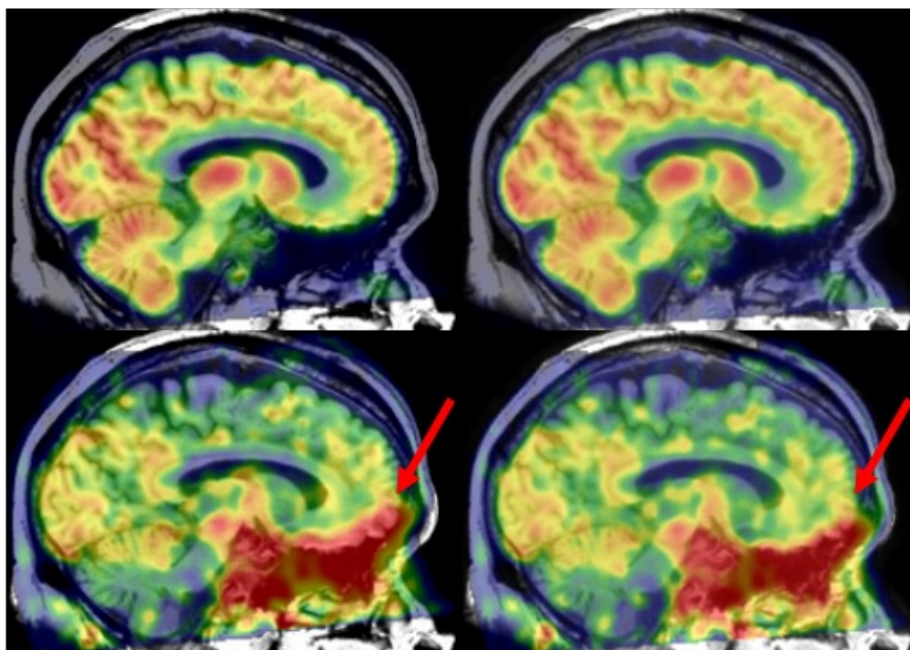
Generate Parametric
Image(s)

Standard SPM
Realignment

Modified SPM
Realignment

First Frame
0-2 min

Last Frame
85-90 min



[¹⁸F]MK-6240

MR-Guided PET Image Processing

Process MRI
(ROI Parcellation)

Smooth/De-noise

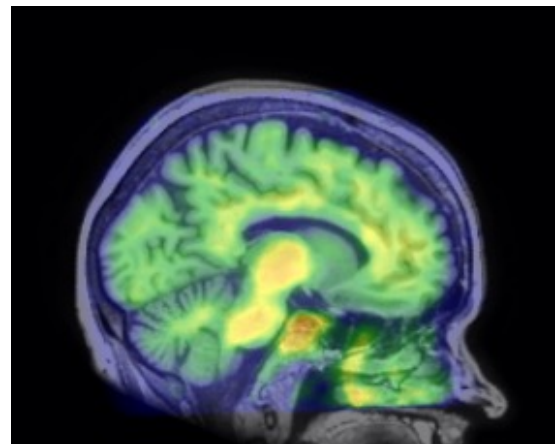
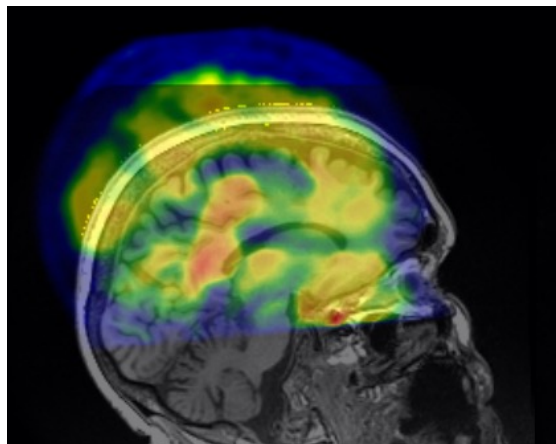
Interframe Alignment
(i.e., motion correction)

Co-Registration to MRI

Extract Reference
Region TAC

Generate Parametric
Image(s)

Intermodal Registration (PET to T1-w MRI)
Unregistered Registered



Reference Image: MRI

Source Image: SUM PET

Process MRI
(ROI Parcellation)

Smooth/De-noise

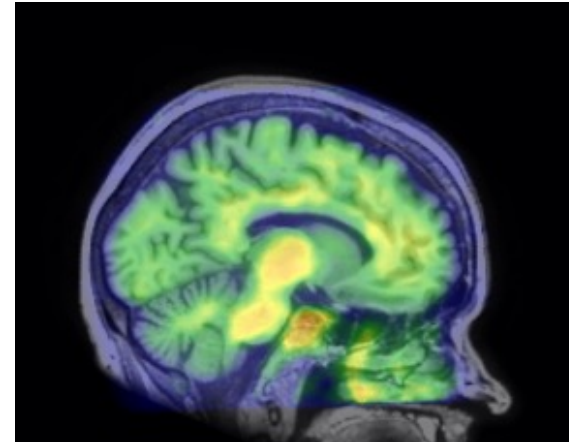
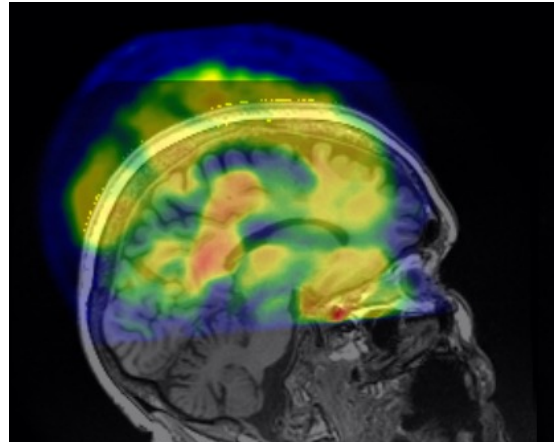
Interframe Alignment
(i.e., motion correction)

Co-Registration to MRI

Extract Reference
Region TAC

Generate Parametric
Image(s)

Intermodal Registration (PET to T1-w MRI)
Unregistered Registered



- Can apply transformation to the PET image using header OR
- Can **reslice** the registered PET image to match voxel-voxel with MRI (requires interpolation but enables extracting ROI-level data)

MR-Guided PET Image Processing

Process MRI
(ROI Parcellation)

Smooth/De-noise

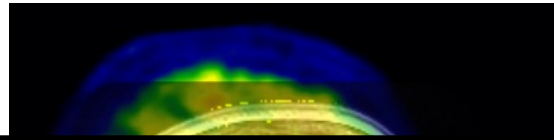
Interframe Alignment
(i.e., motion correction)

Co-Registration to MRI

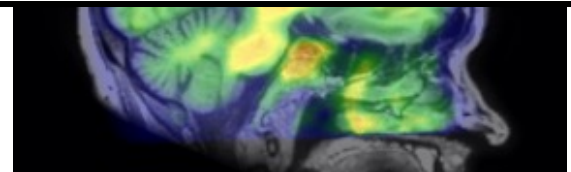
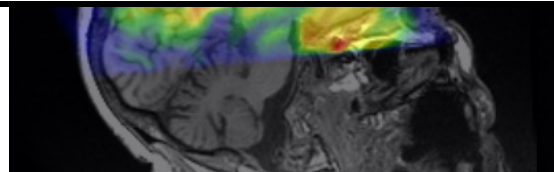
Extract Reference
Region TAC

Generate Parametric
Image(s)

Intermodal Registration (PET to T1-w MRI)
Unregistered Registered



See Previous Basics of Neuroimaging Data Structure and
Formats presented by Dr. Ludovca Griffanti



- Can apply transformation to the PET image using header OR
- Can **reslice** the registered PET image to match voxel-voxel with MRI (requires interpolation but enables extracting ROI-level data)

MR-Guided PET Image Processing

Process MRI
(ROI Parcellation)

Smooth/De-noise

Interframe Alignment
(i.e., motion correction)

Co-Registration to MRI

Extract Reference
Region TAC

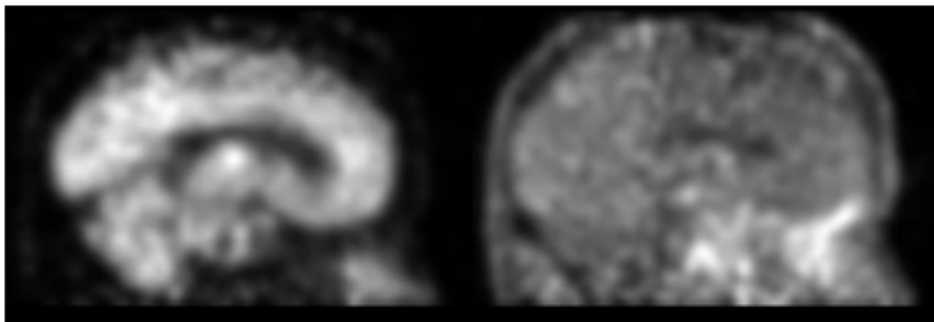
Generate Parametric
Image(s)

Summing Different Frames Can Improve Registration

Early Frame
0-2 min

Late Frame
85-90 min

MK-6240



GM TPM



Process MRI
(ROI Parcellation)

Smooth/De-noise

Interframe Alignment
(i.e., motion correction)

Co-Registration to MRI

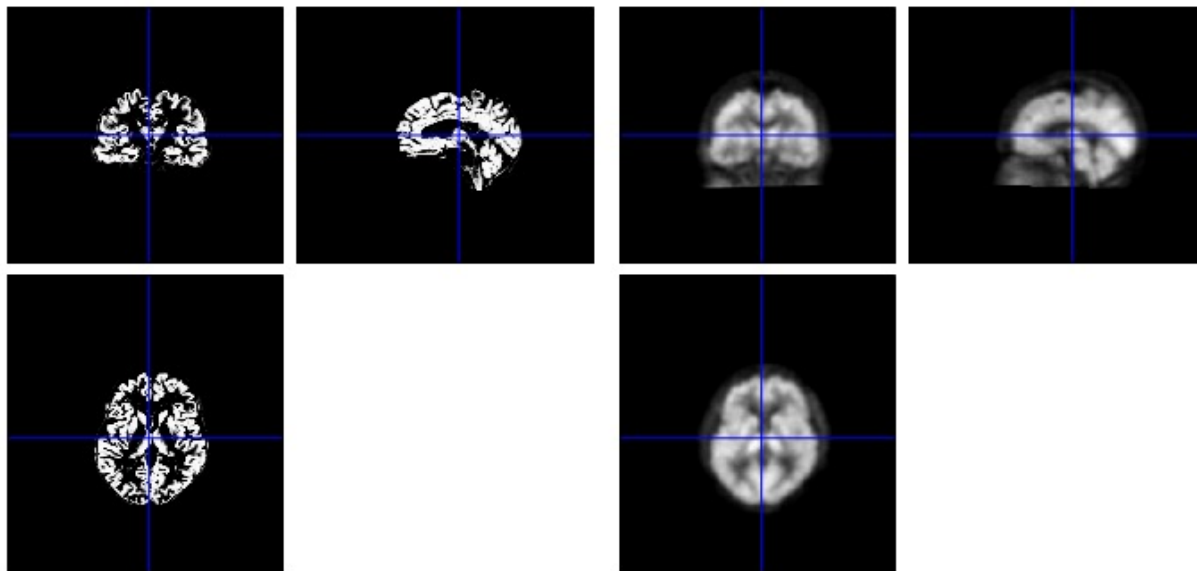
Extract Reference
Region TAC

Generate Parametric
Image(s)

Summing Different Frames Can Improve Registration

GM TPM

MK-6240 SUM 0-10 minutes



- More mutual information between early frame-data and GM TPM compared to late-frame data

MR-Guided PET Image Processing

Process MRI
(ROI Parcellation)

Smooth/De-noise

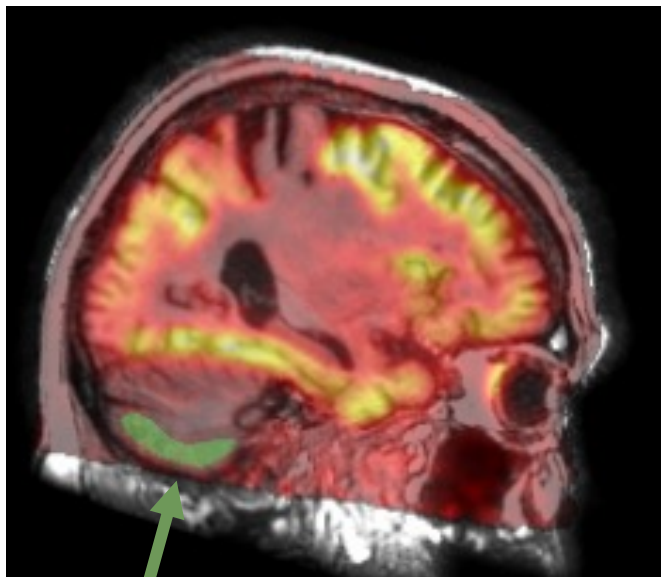
Interframe Alignment
(i.e., motion correction)

Co-Registration to MRI

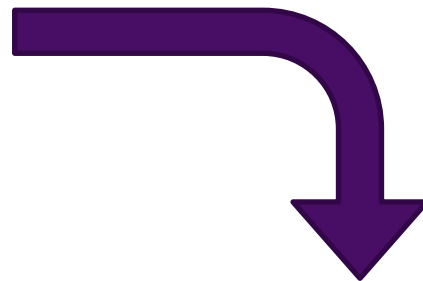
Extract Reference
Region TAC

Generate Parametric
Image(s)

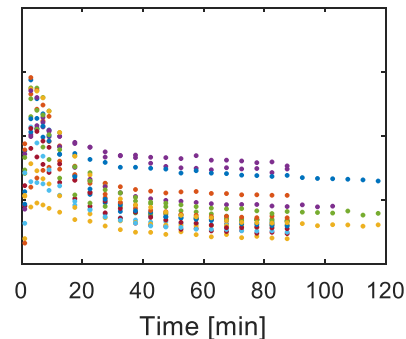
Coregistered PET to MRI



Reference Region
VOI



Time-Activity Curves



Process MRI
(ROI Parcellation)

Smooth/De-noise

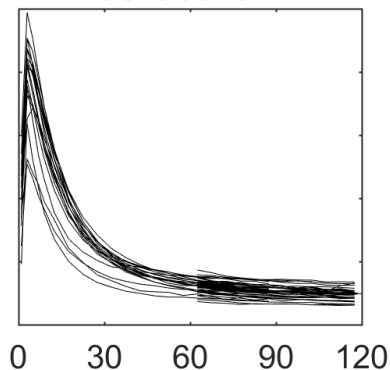
Interframe Alignment
(i.e., motion correction)

Co-Registration to MRI

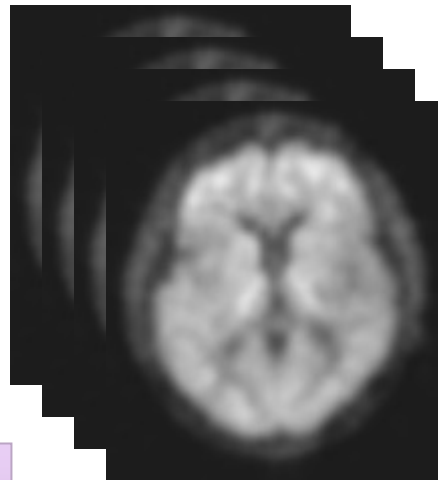
Extract Reference
Region TAC

Generate Parametric
Image(s)

Cerebellum



4D PET



$$\frac{\int_0^T C(t) dt}{C(T)} = \text{DVR} \left[\frac{\int_0^T C_{ref}(t) dt + C_{ref}(T)/\overline{k_2}}{C(T)} \right] + \text{int}$$

MR-Guided PET Image Processing

Process MRI
(ROI Parcellation)

Smooth/De-noise

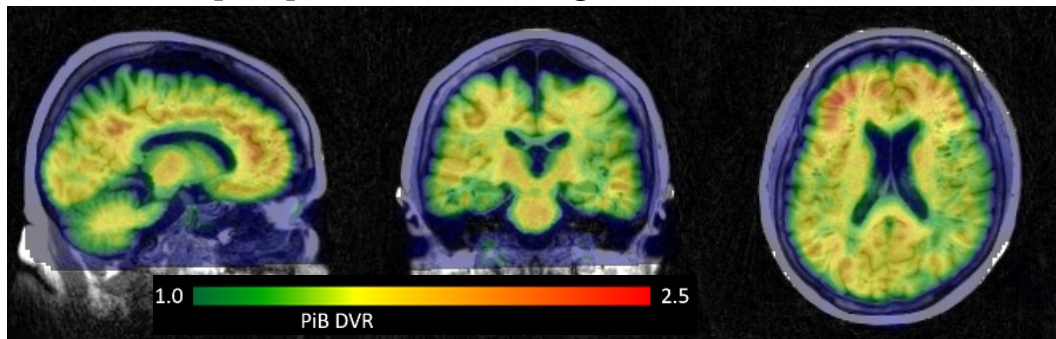
Interframe Alignment
(i.e., motion correction)

Co-Registration to MRI

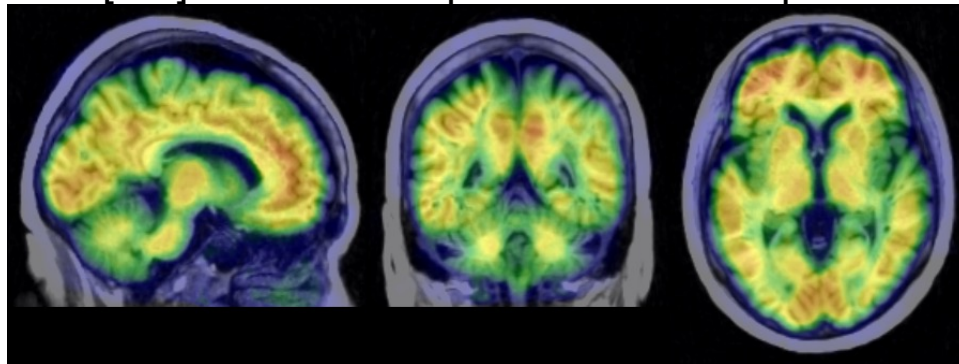
Extract Reference
Region TAC

Generate Parametric
Image(s)

[¹¹C]PiB DVR Coregistered to MRI



[¹¹C]PiB DVR Warped to MNI-152 Space



Apply MRI deformation
field from T1-w spatial
normalization

Process MRI
(ROI Parcellation)

Smooth/De-noise

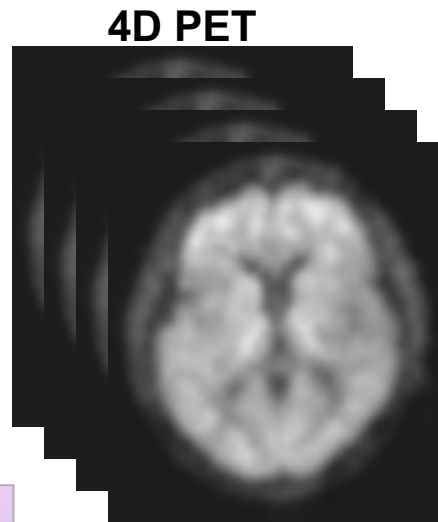
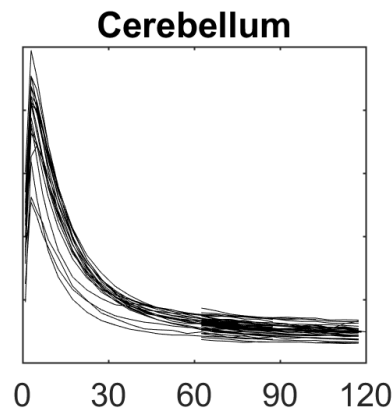
Interframe Alignment
(i.e., motion correction)

Co-Registration to MRI

Extract Reference
Region TAC

Generate Parametric
Image(s)

Can generate other Parametric Images also (e.g., R_1)



Simplified Reference Tissue Method (SRTM)

$$C(T) = \mathbf{R}_1 C_{ref}(t) + \left\{ k_2^{ref} - \frac{R_1 k_2^{ref}}{1 + BP} \right\} C_{ref}(t) \otimes e^{-\left[\frac{k_2^{ref}}{(1+BP)} + \lambda \right] t}$$

Process MRI
(ROI Parcellation)

Smooth/De-noise

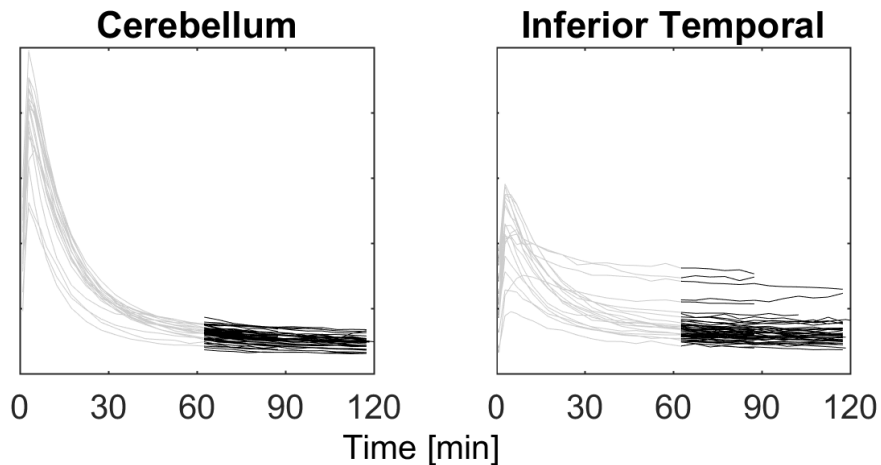
Interframe Alignment
(i.e., motion correction)

Co-Registration to MRI

Extract Reference
Region TAC

Generate Parametric
Image(s)

For **SUVR** quantification, we only have late-frame data



$$\frac{\int_0^T C(t) dt}{C(T)} = DVR \left[\frac{\int_0^T C_{ref}(t) dt + C_{ref}(T)/\bar{k}_2}{C(T)} \right] + int$$

Process MRI
(ROI Parcellation)

Smooth/De-noise

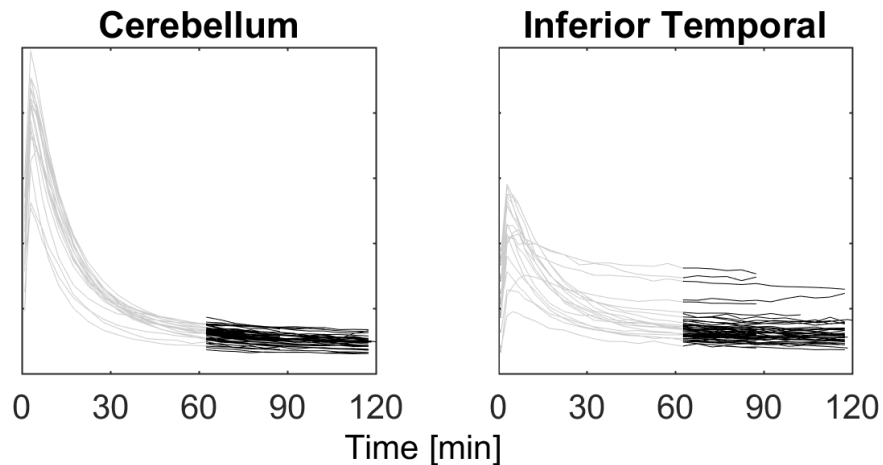
Interframe Alignment
(i.e., motion correction)

Co-Registration to MRI

Extract Reference
Region TAC

Generate Parametric
Image(s)

For **SUVR** quantification, we only have late-frame data



$$SUVR = \frac{C(\Delta t)}{C_{ref}(\Delta t)}$$

Process MRI
(ROI Parcellation)

Smooth/De-noise

Interframe Alignment
(i.e., motion correction)

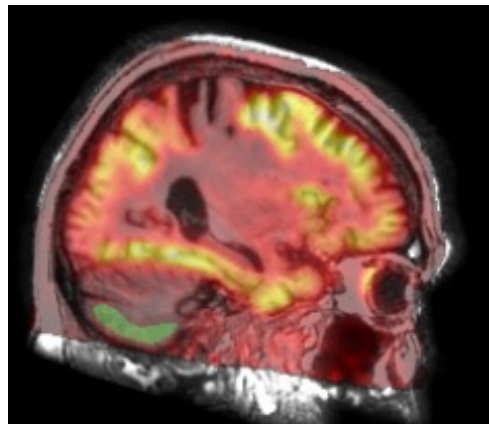
Co-Registration to MRI

Extract Reference
Region TAC

Generate Parametric
Image(s)

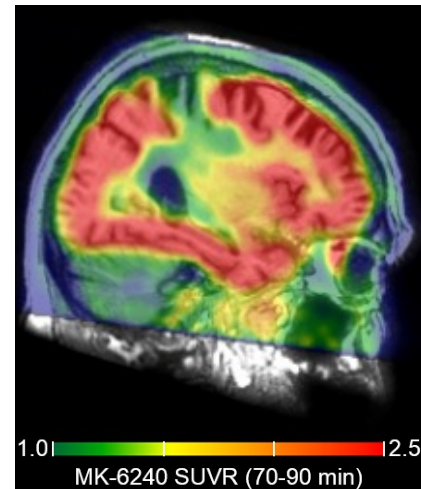
Create SUVR image by dividing entire image by mean activity concentration in the reference region

Coregistered Summed PET



$$\frac{I_{SUM}}{\text{mean}(I_{SUM,ref})}$$

SUVR Image



MR-Guided PET Image Processing

Process MRI
(ROI Parcellation)

Smooth/De-noise

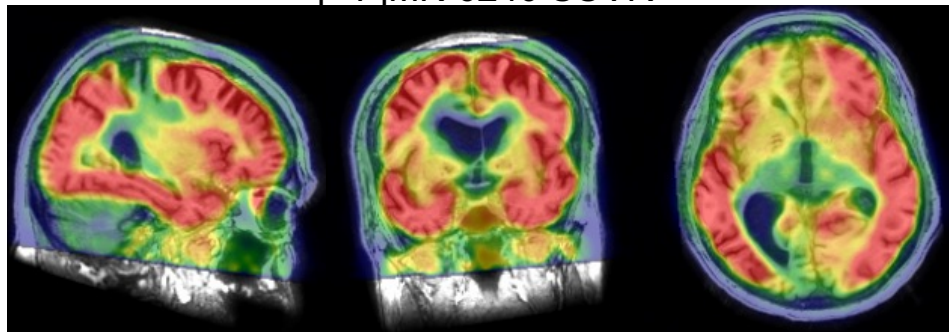
Interframe Alignment
(i.e., motion correction)

Co-Registration to MRI

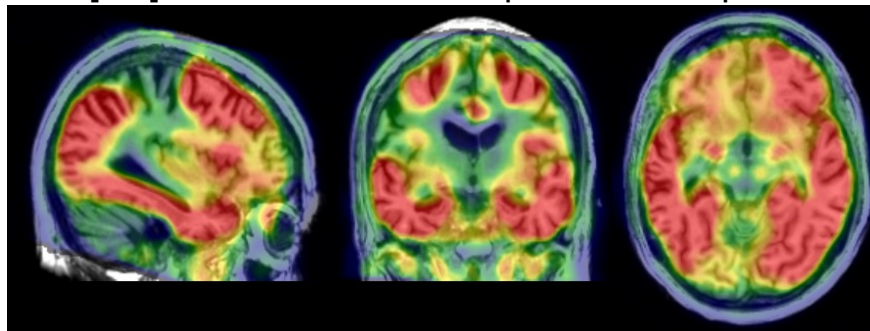
Extract Reference
Region TAC

Generate Parametric
Image(s)

[¹⁸F]MK-6240 SUVR



[¹⁸F]MK-6240 SUVR Warped to MNI Space



Apply MRI deformation
field from T1-w spatial
normalization

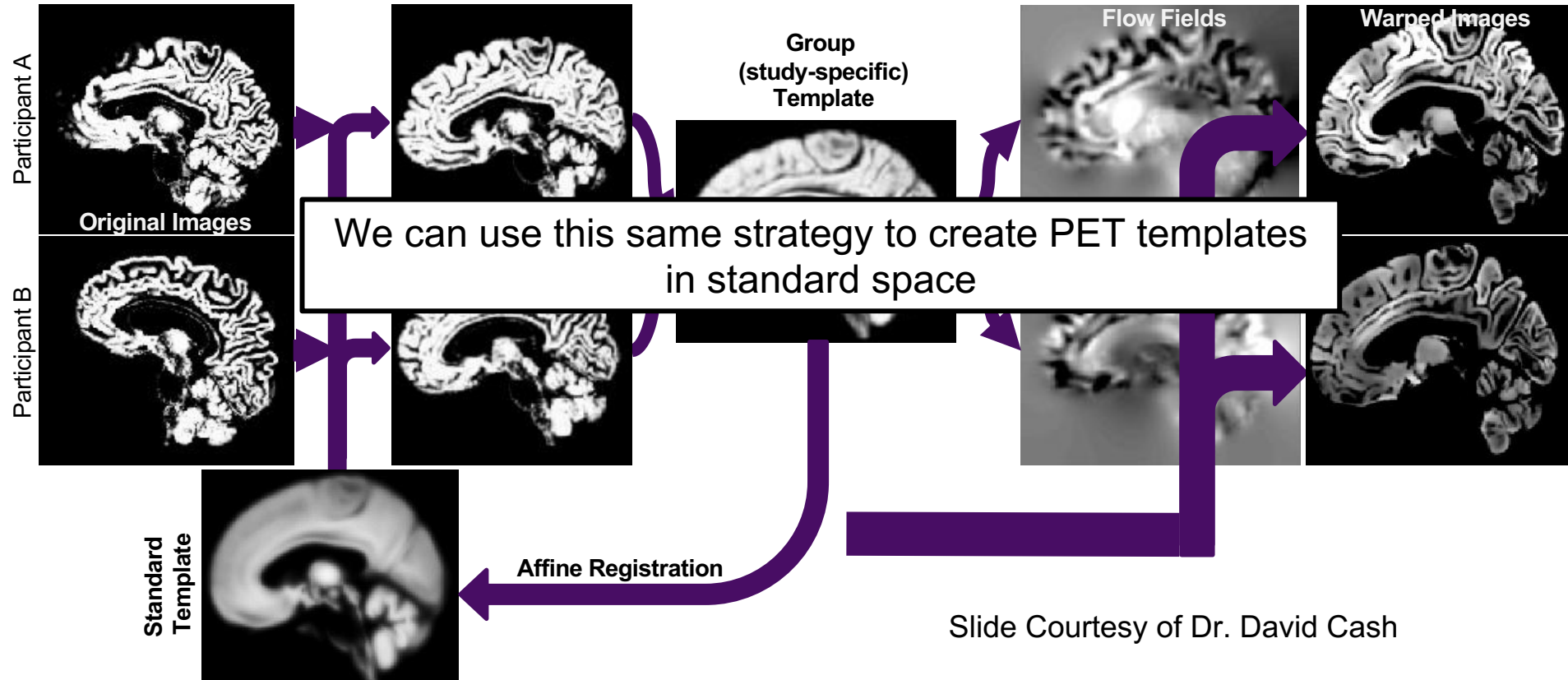
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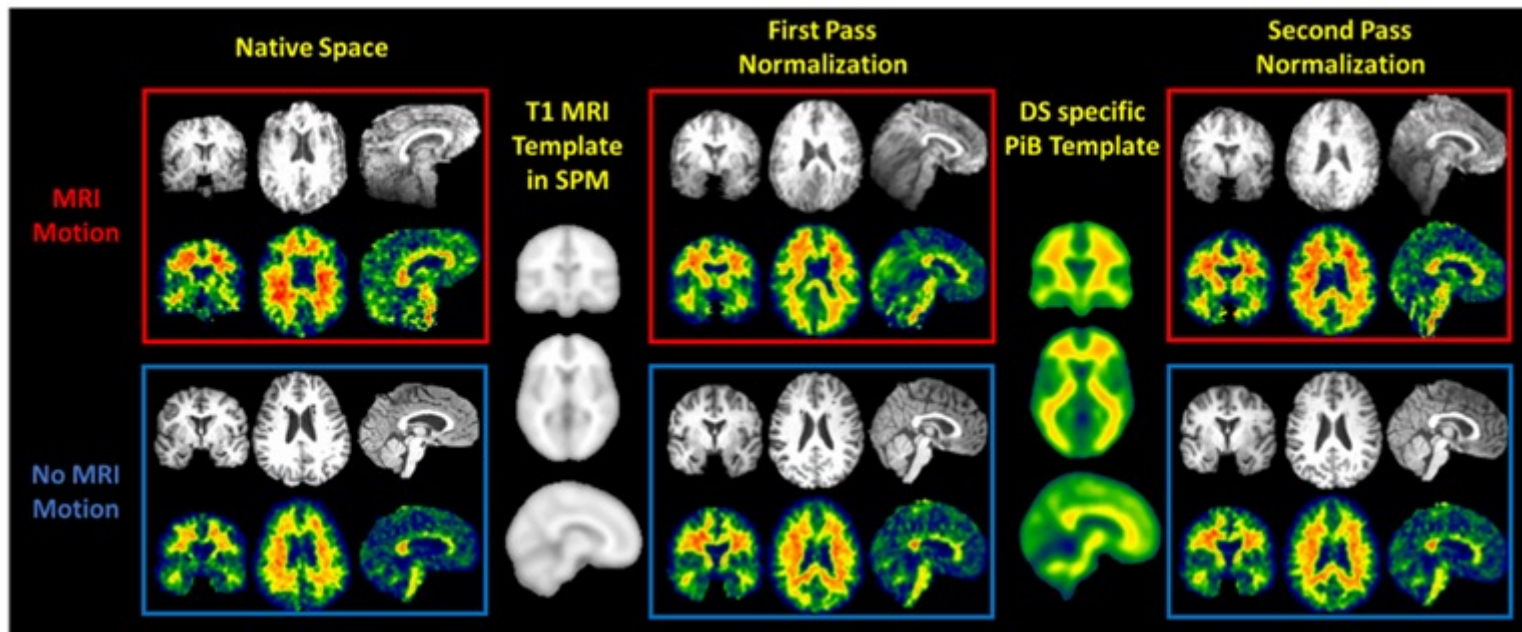
AAIC>23

PET Image Processing

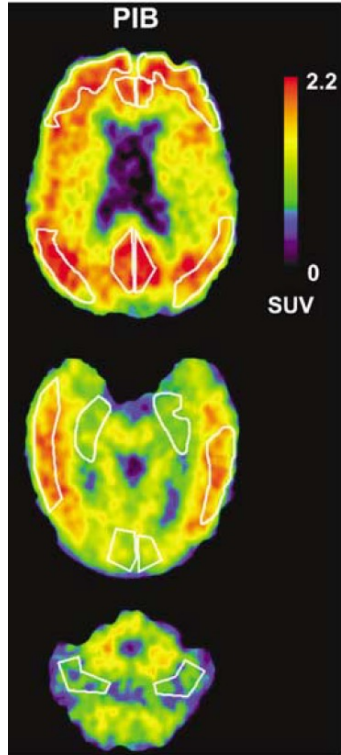
- MR-Guided Image Processing
- **PET only Image Processing**
- Other Considerations

Creating a study-specific template to reduce errors in spatial normalisation





Lao, et al., Brain Imaging and Behavior, 2019



ROIs can also be Hand drawn on
SUM or SUV Images

- Create SUM PET image
- Manually draw ROIs on image
- Extract TACs or regional mean
- Generate parametric image

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PET Image Processing

- MR-Guided Image Processing
- PET only Image Processing
- **Other Considerations**

- Reconstruction parameters (corrections for deadtime, scatter, attenuation, decay, etc.,)
- Standardization across tracers, sites, acquisition protocols, etc.,
- Partial Volume Effects
- Reference region selection
- Off-target binding
- Brain-penetrable radiometabolites

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POP QUIZ!

Which of the following statements best describes PET imaging?:

- a) High spatial resolution, high molecular specificity
- b) Low spatial resolution, high molecular specificity
- c) High spatial resolution, low molecular specificity
- d) Low spatial resolution, low molecular specificity

Which of the following statements best describes PET imaging?:

- a) High spatial resolution, high molecular specificity
- b) Low spatial resolution, high molecular specificity**
- c) High spatial resolution, low molecular specificity
- d) Low spatial resolution, low molecular specificity

The signal we detect with PET imaging is:

- a) Single gamma photons
- b) Beta particles
- c) X-rays
- d) Coincident gamma photons

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PET radiotracers for amyloid and tau mostly reflect:

- a) Soluble protein fragments
- b) Transient pathological changes in beta-amyloid and tau
- c) Insoluble protein aggregates
- d) None of the above

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Registering PET to MRI provides:

- a) Less noisy images
- b) Anatomical reference and regions of interest
- c) The underlying radiotracer distribution at higher resolution
- d) Regional radiotracer perfusion information

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A SUM PET image is:

- a) A time-weighted average of all or some PET frames
- b) A quantitative measure of binding potential
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- d) A quantitative measure of perfusion

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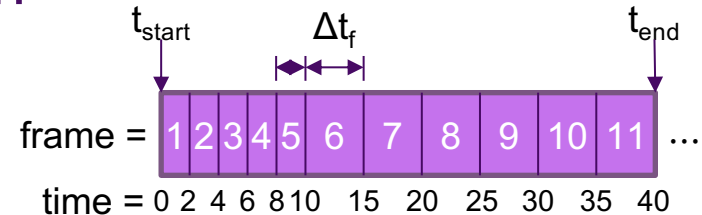
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List Mode

$$\sum_{i=t_{start}}^{t_{end}} \frac{counts_i}{t_i}$$

Reconstructed Image Frames

$$\frac{\sum_{f=frame_{start}}^{frame_{end}} C(t)_f \times \Delta t_f}{\sum_{f=frame_{start}}^{frame_{end}} \Delta t_f}$$



THANK YOU!



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Past Webinars in this Series:

- Basics of Neuroimaging: Data structure and formats by Ludovica Griffanti
- Basics of Neuroimaging: Structural MRI by David Cash

On demand at <https://training.alz.org/research-webinars>

Next up:

- Basics of Neuroimaging: Diffusion-Weighted Imaging (DWI) by Alexa Pichet Binette
21 April, 2023; 9AM – 10AM CT
- Basics of Neuroimaging: Functional Magnetic Resonance Imaging (fMRI) by Luigi Lorenzini
26 April, 2023; 10AM – 11AM CT

GETTING STARTED WITH NEUROIMAGING WORKSHOP Friday, July 14 8:00-12:00 Amsterdam