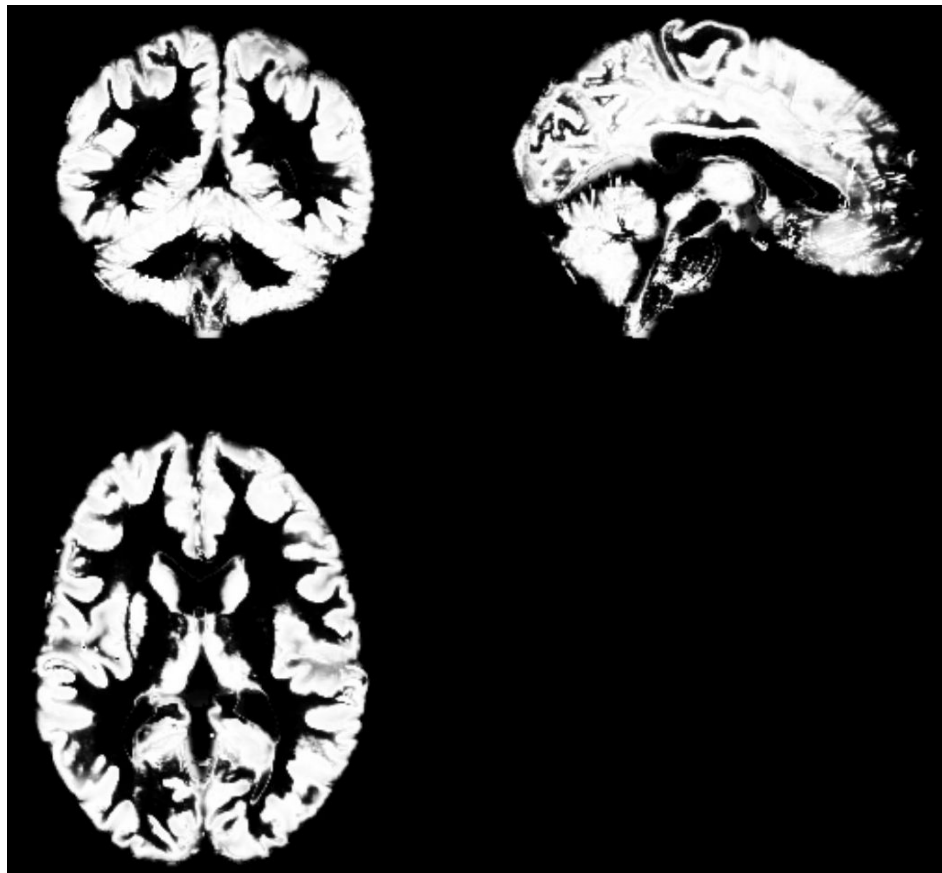


The Transition from Physical to Abstract Taste: An fMRI Study on Sweet, Sour, and Salty

by Tran Cao Bang Trinh 陳高冰貞 (Annamarie)
Graduate Institute of Linguistics, NTU (TW)
1st year PhD Student

Motivation



Segmented brain
Gray matter
Sub-007
1st study

Introduction:

This ice cream tastes **sweet!** (Physical)

You're so **sweet!** (Abstract)

Introduction:

- **Embodiment theory** posits that abstract thought is grounded in concrete, sensorimotor experiences.
- **Taste**, while fundamentally a **physical** sensation, is frequently extended into the **abstract** domain (e.g., "sweet success," "bitter truth").
- This conceptual overlap suggests that abstract uses of taste may recruit **similar neural mechanisms** as literal, physical taste experiences.

Hypothesis:

- **Abstract** taste expressions will partially **overlap** with **physical** taste processing regions (e.g., insula, primary gustatory cortex).
- This overlap will be spatially reduced, and abstract taste will additionally recruit regions associated with semantic and conceptual processing.
- Connectivity between abstract-related areas and primary taste regions will reflect grounding in physical experience.

Participant

Best Pair (female)

- **Dataset 1:** sub-007 (68 kg, 24, F)
- **Dataset 2:** sub-072 (65.5 kg, 26, F)
 - *Same gender, age difference = 2 years, weight difference = 2.5 kg.*

Methodology

1. 1st level analysis: Visual inspection across both group by best pair.
 - Choosing a comparable pair from both group.
 - Preprocessing data > 1st level analysis (SPM12).
 - Visual inspection.
2. 2nd level analysis: Comparison across both group.
 - Adding comparable pairs (female, male, different ages).
 - Including all participants.

Source:

Taste Quality Representation in the Human Brain (2020)

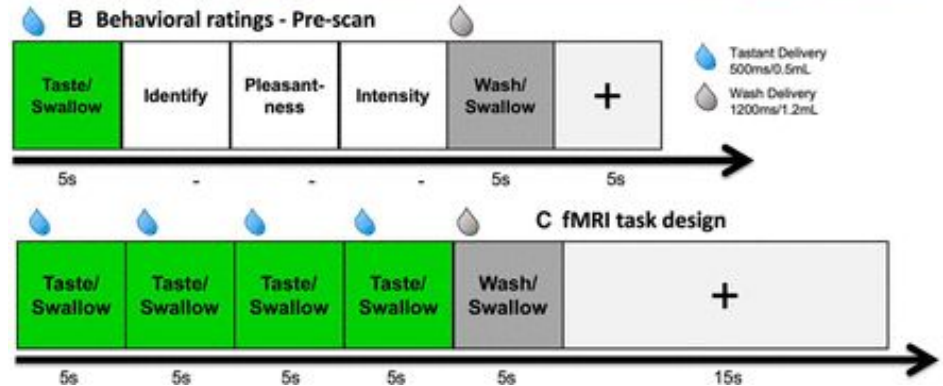
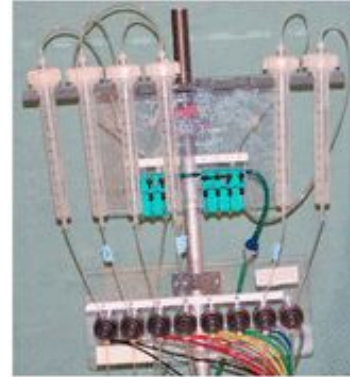
A common neural code for representing imagined and inferred tastes (2023)

1st study:

onset duration trial_type

5	20	sour
25	5	wash
45	20	neutral
65	5	wash
85	20	salty
105	5	wash
125	20	sweet
145	5	wash
165	20	sour
185	5	wash
205	20	salty
225	5	wash
245	20	neutral
265	5	wash
285	20	sweet
305	5	wash

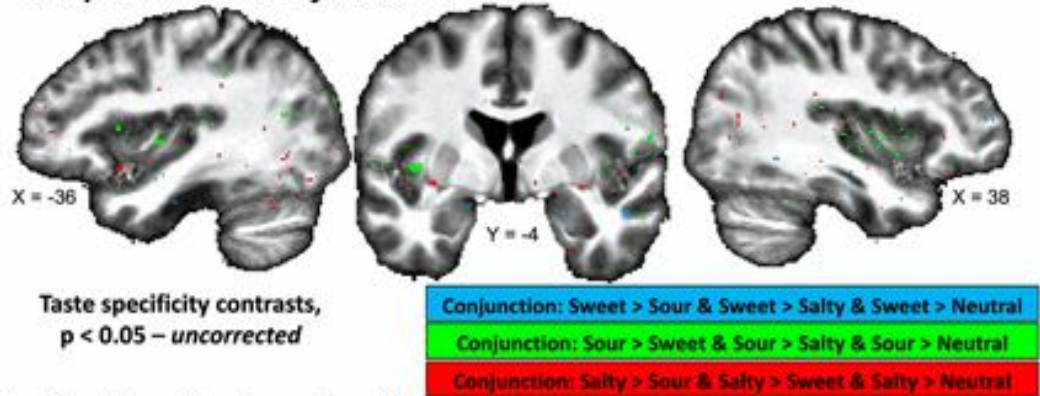
*8 runs



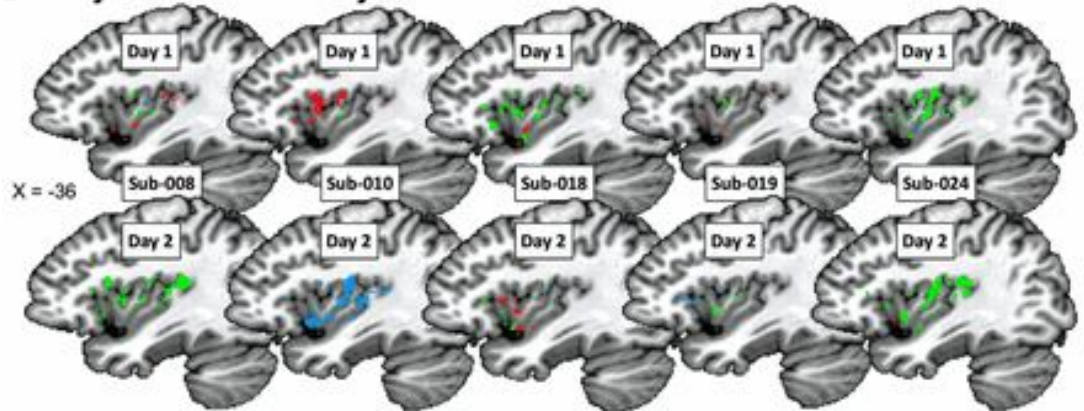
1st study:

The brain **does not** have strongly segregated, stable topographic maps for **basic tastes** (sweet, salty, sour) at the group level, and even **individual subjects** showed inconsistent patterns across sessions.

A Group level taste conjunctions

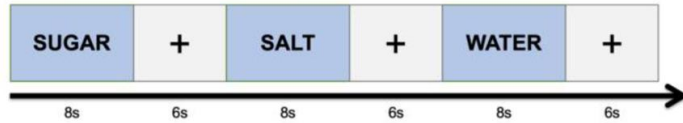


B Subject-level taste conjunctions

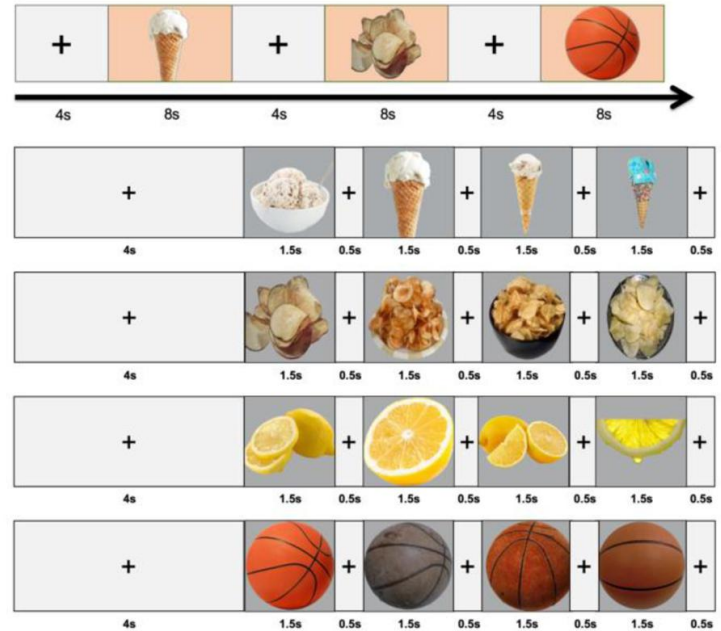


2nd study:

A Taste Imagining fMRI Task



B Food Pictures fMRI Task



2nd study:

Tatseimagine- Run1/4

Foodpicture-Run 1/4

onset	duration	trial_type
--------------	-----------------	-------------------

6000	8000	water
------	------	-------

20000	8000	lemon
-------	------	-------

34000	8000	sugar
-------	------	-------

48000	8000	lemon
-------	------	-------

62000	8000	water
-------	------	-------

76000	8000	salt
-------	------	------

90000	8000	sugar
-------	------	-------

onset	duration	trial_type
--------------	-----------------	-------------------

4000	8000	salty
------	------	-------

16000	8000	sour
-------	------	------

28000	8000	sweet
-------	------	-------

40000	8000	sour
-------	------	------

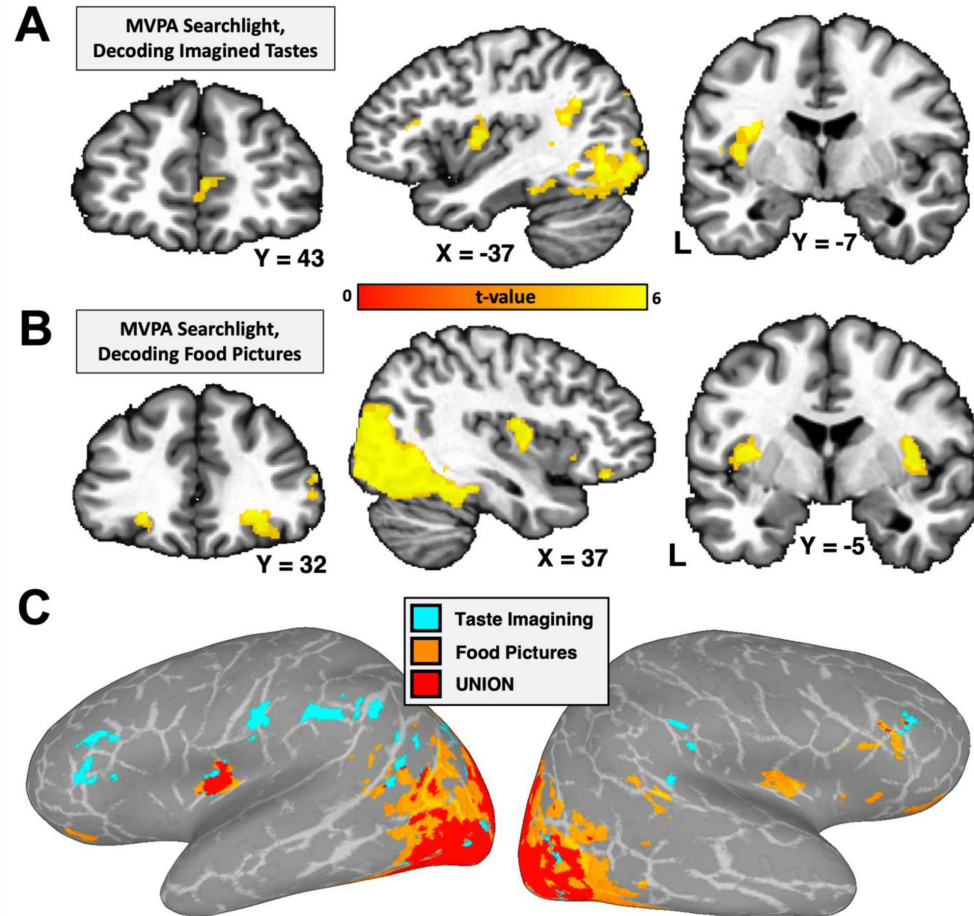
52000	8000	object
-------	------	--------

64000	8000	sour
-------	------	------

76000	8000	object
-------	------	--------

2nd study:

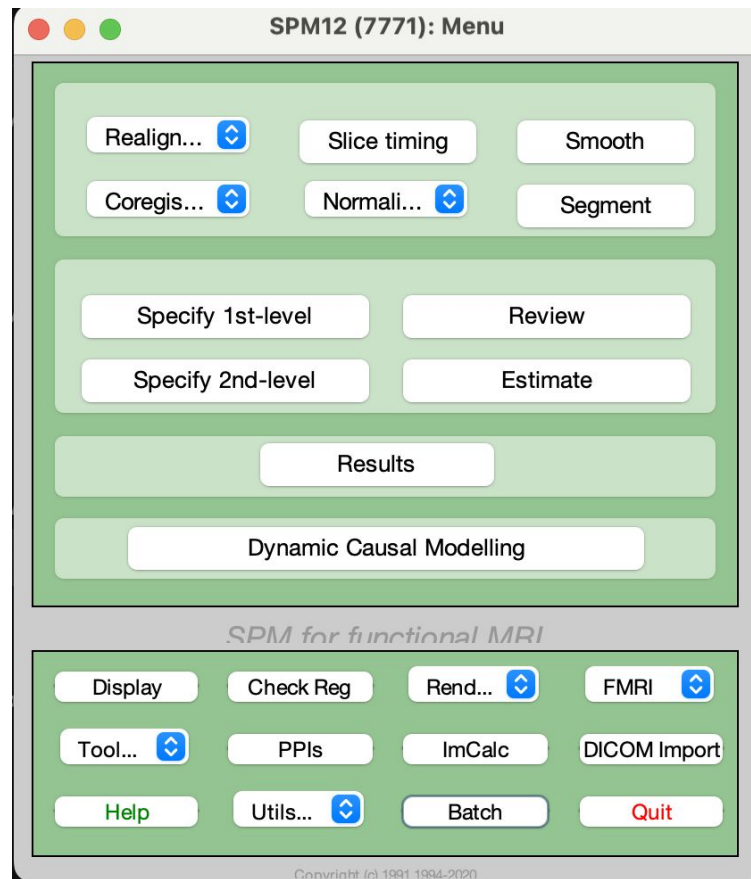
- No brain region analysis for specific taste
- Inclusion of objects may affect the results for Food picture tasks.



Pipeline

Preprocessing pipeline: SPM

- Check Reg (Motion Correction)
- Realign & Reslice
- Slice timing
- Coregister T1 EPI
- Segmentation



Data downloading

Download with DataLad

Public datasets can be downloaded with [DataLad](#) or [Git Annex](#) from [GitHub](#).

```
datalad install https://github.com/OpenNeuroDatasets/ds002995.git
```

Check out [getting started with DataLad](#) for more on how to use this download method.

Download with a shell script

A script is available to download with only curl. This may be useful if your download environment makes it difficult to install DataLad or Node.js.

[Download shell script](#)

Datalab

```
[PATH ...]
(base) MacBook-Pro-de-Annamarie:~ annamarie$ cd /Users/annamarie/ds004312
[(base) MacBook-Pro-de-Annamarie:ds004312 annamarie$ datalad get sub-072
]
[get(ok): sub-072/ses-01/anat/sub-072_ses-01_T1w.nii.gz (file) [from s3-PUBLIC...]]

get(ok): sub-072/ses-01/fmap/sub-072_ses-01_dir-PA_epi.nii.gz (file) [from s3-PUBLIC...]
get(ok): sub-072/ses-01/func/sub-072_ses-01_task-foodpicture_run-01_bold.nii.gz (file) [from s3-PUBLIC...]
get(ok): sub-072/ses-01/func/sub-072_ses-01_task-foodpicture_run-02_bold.nii.gz (file) [from s3-PUBLIC...]
get(ok): sub-072/ses-01/func/sub-072_ses-01_task-foodpicture_run-03_bold.nii.gz (file) [from s3-PUBLIC...]
get(ok): sub-072/ses-01/func/sub-072_ses-01_task-foodpicture_run-04_bold.nii.gz (file) [from s3-PUBLIC...]
get(ok): sub-072/ses-01/func/sub-072_ses-01_task-tasteimagine_run-01_bold.nii.gz (file) [from s3-PUBLIC...]
get(ok): sub-072/ses-01/func/sub-072_ses-01_task-tasteimagine_run-02_bold.nii.gz (file) [from s3-PUBLIC...]
get(ok): sub-072/ses-01/func/sub-072_ses-01_task-tasteimagine_run-03_bold.nii.gz (file) [from s3-PUBLIC...]
get(ok): sub-072/ses-01/func/sub-072_ses-01_task-tasteimagine_run-04_bold.nii.gz (file) [from s3-PUBLIC...]
get(ok): sub-072 (directory)
action summary:
  get (ok: 11)
(base) MacBook-Pro-de-Annamarie:ds004312 annamarie$ █
```

=> amex-encoded file

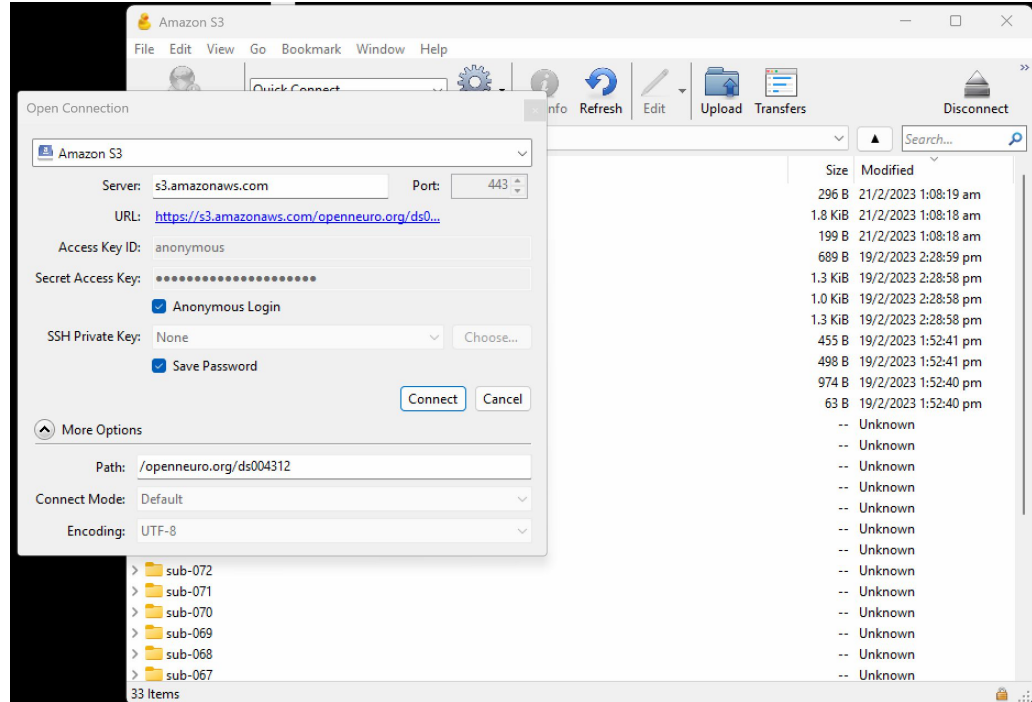
Shell script

```
1 #!/bin/sh
2 curl --create-dirs https://s3.amazonaws.com/openneuro.org/ds002995/CHANGES?versionId=kctH4K.Ji68clv8NJBhw4_LOWJIUBweh -o ds002995-1.0.1/CHANGES
3 curl --create-dirs https://s3.amazonaws.com/openneuro.org/ds002995/README?versionId=pd6tB8NAJWjmbvrA2mVcby5n_zL_Zk2n -o ds002995-1.0.1/README
4 curl --create-dirs https://s3.amazonaws.com/openneuro.org/ds002995/dataset_description.json?versionId=g6YfEuE4BTq04nhZlzxMjZBj2U9.t2af -o ds002995-1.0.1/dataset_description.json
5 curl --create-dirs https://s3.amazonaws.com/openneuro.org/ds002995/.README?versionId=_I9pPUBNEubaYHCbv98wWPjjqQpN2RfL -o ds002995-1.0.1/.README
6 curl --create-dirs https://s3.amazonaws.com/openneuro.org/ds002995/T1w.json?versionId=yb4FTs.yKyIKOr1dUZY0uJd.UJhvg3r -o ds002995-1.0.1/T1w.json
7 curl --create-dirs https://s3.amazonaws.com/openneuro.org/ds002995/dir-PA_epi.json?versionId=yf0eZoTs3fpU_izMDOcLmLZHtnTdLptG -o ds002995-1.0.1/dir-PA_epi.json
8 curl --create-dirs https://s3.amazonaws.com/openneuro.org/ds002995/participants.json?versionId=dPwt_WzP9uTOxZv01q3m022Zhp9FQiW -o ds002995-1.0.1/participants.json
9 curl --create-dirs https://s3.amazonaws.com/openneuro.org/ds002995/participants.tsv?versionId=WoLgUvkZh2ZZ8wxMadMFhnyxr0VE6r_A -o ds002995-1.0.1/participants.tsv
10 curl --create-dirs https://s3.amazonaws.com/openneuro.org/ds002995/task-tastemap_bold.json?versionId=VEbvZFCLNdPhsnbAsTvXB0qJaJisU5mG -o ds002995-1.0.1/task-tastemap_bold.json
11 curl --create-dirs https://s3.amazonaws.com/openneuro.org/ds002995/task-tastemap_events.json?versionId=Qe1IbfxyFhWxVXQq9ZJsuQBfUprJBx7l -o ds002995-1.0.1/task-tastemap_events.json
12 curl --create-dirs https://s3.amazonaws.com/openneuro.org/ds002995/sub-007/ses-01/anat/sub-007_ses-01_T1w.nii.gz?versionId=HT7UVYG2yxqAyQjPuWM9XbIHbDFyybV3 -o
    ds002995-1.0.1/sub-007/ses-01/anat/sub-007_ses-01_T1w.nii.gz
13 curl --create-dirs https://s3.amazonaws.com/openneuro.org/ds002995/sub-007/ses-01/fmap/sub-007_ses-01_dir-PA_epi.nii.gz?versionId=d7l7IrCwNZSs5W.ZuiA2LJEX07dTJQ6R -o
    ds002995-1.0.1/sub-007/ses-01/fmap/sub-007_ses-01_dir-PA_epi.nii.gz
14 curl --create-dirs https://s3.amazonaws.com/openneuro.org/ds002995/sub-007/ses-01/func/.sub-007_ses-01_task-tastemap_run-01_events.tsv?versionId=zyv_x2cosDkby4eos4zok.Synu8ShmcF
    -o ds002995-1.0.1/sub-007/ses-01/func/.sub-007_ses-01_task-tastemap_run-01_events.tsv
15 curl --create-dirs https://s3.amazonaws.com/openneuro.org/ds002995/sub-007/ses-01/func/.sub-007_ses-01_task-tastemap_run-02_events.tsv?versionId=N8eVZhskDmKcuYJt0J8f1kRMB95jR0z
    -o ds002995-1.0.1/sub-007/ses-01/func/.sub-007_ses-01_task-tastemap_run-02_events.tsv
16 curl --create-dirs https://s3.amazonaws.com/openneuro.org/ds002995/sub-007/ses-01/func/.sub-007_ses-01_task-tastemap_run-03_events.tsv?versionId=KKncroUKJ4CbBZ4e0isBIcChFSBvd7aM
    -o ds002995-1.0.1/sub-007/ses-01/func/.sub-007_ses-01_task-tastemap_run-03_events.tsv
```

=> complicated

Cyberduck > Amazon S3

=> Path spec



Slice-timing

=> Check README, json file.

```
"Columns": 1440,  
"PixelSpacing": "1.2000000476837 1.2000000476837",  
"SliceTiming": [  
    0.000,  
    1.845,  
    1.183,  
    0.517,  
    2.362,  
    1.697,
```

Matlab > Slice-timing info

```
>> % Step 1: Define your SliceTiming (from your JSON)
SliceTiming = [ ...
    0.000, 1.845, 1.183, 0.517, 2.362, 1.697, 1.035, 0.370, ...
    2.215, 1.550, 0.887, 0.222, 2.068, 1.403, 0.738, 0.075, ...
    1.920, 1.255, 0.590, 2.435, 1.773, 1.107, 0.442, 2.287, ...
    1.625, 0.960, 0.295, 2.140, 1.477, 0.813, 0.147, 1.992, ...
    1.330, 0.665, 0.000, 1.845, 1.183, 0.517, 2.362, 1.697, ...
    1.035, 0.370, 2.215, 1.550, 0.887, 0.222, 2.068, 1.403, ...
    0.738, 0.075, 1.920, 1.255, 0.590, 2.435, 1.773, 1.107, ...
    0.442, 2.287, 1.625, 0.960, 0.295, 2.140, 1.477, 0.813, ...
    0.147, 1.992, 1.330, 0.665 ];

% Step 2: Compute slice acquisition order
[~, slice_order] = sort(SliceTiming);

% Step 3: Find reference slice (closest to 1.0s)
[~, ref_slice] = min(abs(SliceTiming - 1.0));

% Step 4: Display results
fprintf('Slice acquisition order:\n');
disp(slice_order);

fprintf('Reference slice (closest to 1s): %d\n', ref_slice);
```

Matlab > Slice-timing info

Slice acquisition order:

Columns 1 through 12

1	35	16	50	31	65	12	46	27	61	8	42
---	----	----	----	----	----	----	----	----	----	---	----

Columns 13 through 24

23	57	4	38	19	53	34	68	15	49	30	64
----	----	---	----	----	----	----	----	----	----	----	----

Columns 25 through 36

11	45	26	60	7	41	22	56	3	37	18	52
----	----	----	----	---	----	----	----	---	----	----	----

Columns 37 through 48

33	67	14	48	29	63	10	44	25	59	6	40
----	----	----	----	----	----	----	----	----	----	---	----

Columns 49 through 60

21	55	2	36	17	51	32	66	13	47	28	62
----	----	---	----	----	----	----	----	----	----	----	----

Columns 61 through 68

9	43	24	58	5	39	20	54
---	----	----	----	---	----	----	----

Reference slice (closest to 1s): 7

Matlab > Slice-timing info

Current Module: Slice Timing	
Help on: Slice Timing	
Data	
. Session	130 files
. Session	130 files
. Session	130 files
. Session	130 files
. Session	130 files
. Session	130 files
. Session	130 files
. Session	130 files
Number of Slices	68
TR	2.5
TA	2.46323529411765
Slice order	1x68 double
Reference Slice	7
Filename Prefix	a

No slice-timing info? Assume or skip ?

Slice Order for Siemens Scanner with 58 Slices

Because:

- **Vendor:** Siemens
- **Acquisition:** 2D EPI
- **Slices:** 58 (even number)
- **No **SliceTiming** provided**

Then according to **Siemens' standard interleaved rule** for **even number of slices**, the slice order is:

```
slice_order = [2:2:58 1:2:57];
```

Reference Slice

SPM needs you to pick a **reference slice** — usually the one acquired **closest to the middle of the TR**, to reduce temporal bias.

Cogister T1 EPI

No T2 image

T1 > EPI

Current Module: Coregister: Estimate

Help on: Coregister: Estimate

Reference Image ...01/func/arsub-007_ses-01_task-tastemap_run-01_bold.nii,1

Source Image .../ds002995/sub-007/ses-01/anat/sub-007_ses-01_T1w.nii,1

Other Images

Estimation Options

. Objective Function

. Separation

. Tolerances

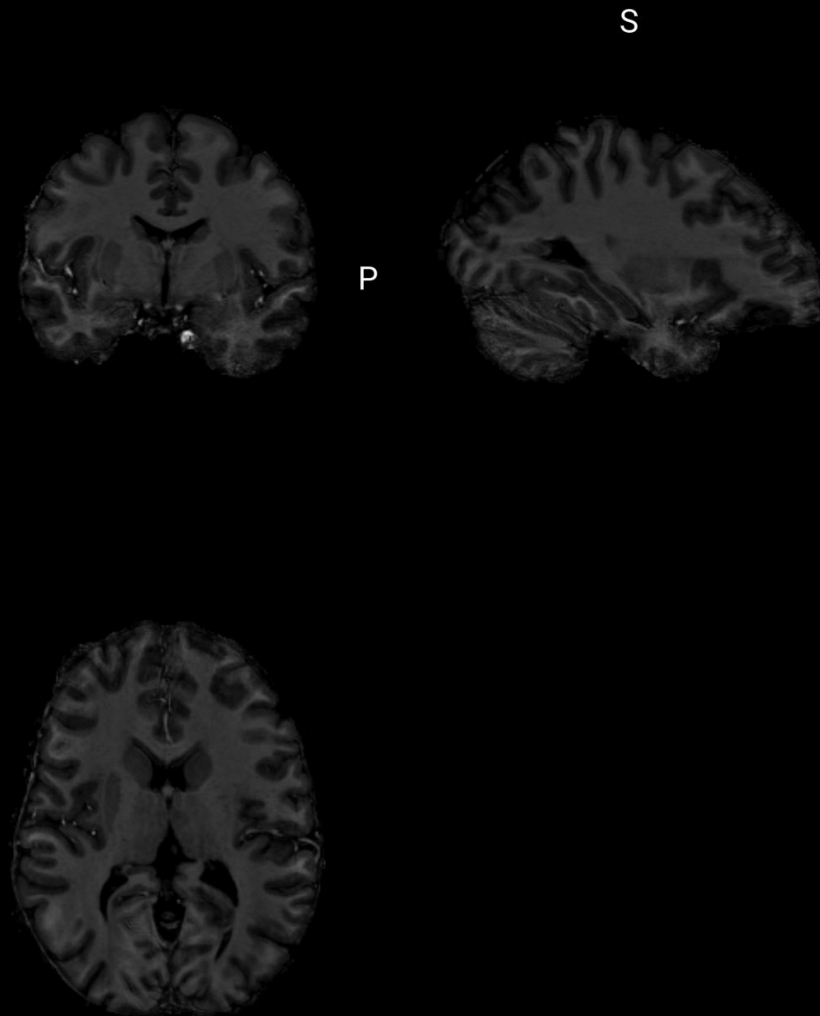
. Histogram Smoothing

Normalised Mutual Information

[4 2]

1x12 double

[7 7]



Missing brain image

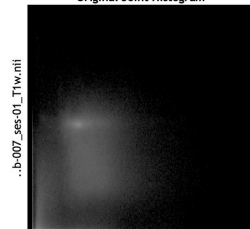
Normalised Mutual Information Coregistration

$$X1 = 0.003 \cdot X - 0.021 \cdot Y + 0.583 \cdot Z + 21.946$$

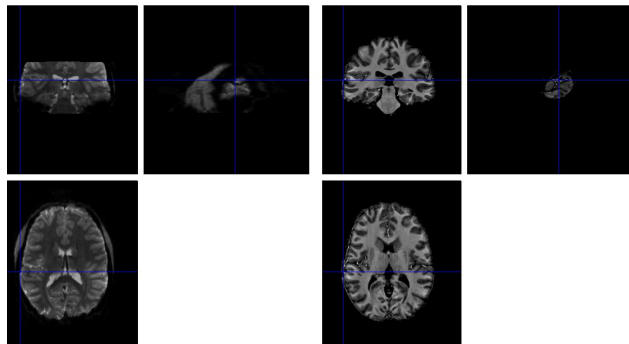
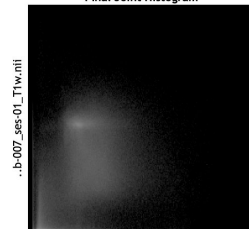
$$Y1 = 0.583 \cdot X - 0.007 \cdot Y - 0.003 \cdot Z - 3.996$$

$$Z1 = -0.007 \cdot X - 0.583 \cdot Y - 0.021 \cdot Z + 132.854$$

Original Joint Histogram



Final Joint Histogram



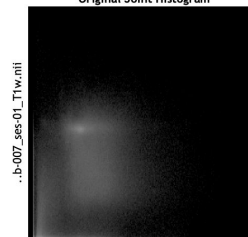
Normalised Mutual Information Coregistration

$$X1 = 0.003 \cdot X - 0.021 \cdot Y + 0.583 \cdot Z + 21.946$$

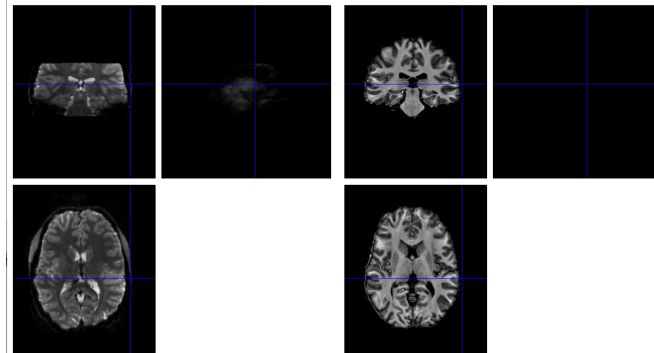
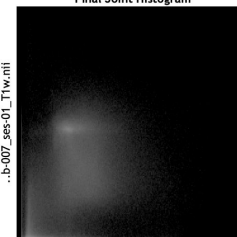
$$Y1 = 0.583 \cdot X - 0.007 \cdot Y - 0.003 \cdot Z - 3.996$$

$$Z1 = -0.007 \cdot X - 0.583 \cdot Y - 0.021 \cdot Z + 132.854$$

Original Joint Histogram



Final Joint Histogram



Missing brain image

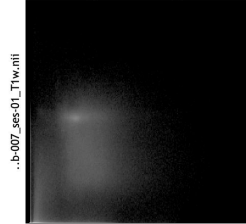
Normalised Mutual Information Coregistration

$$X1 = 0.003^{\circ}X - 0.021^{\circ}Y + 0.583^{\circ}Z + 21.946$$

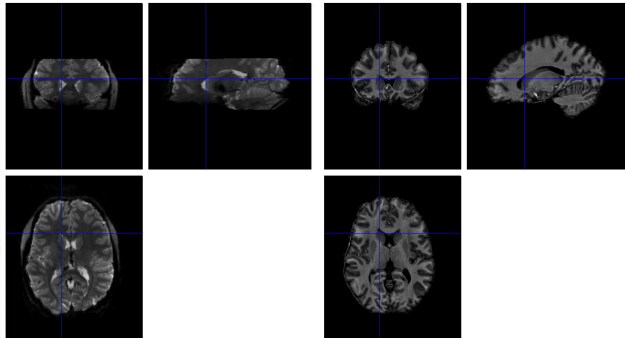
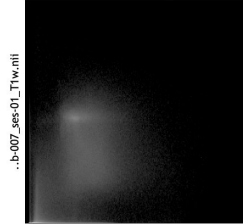
$$Y1 = 0.583^{\circ}X - 0.007^{\circ}Y - 0.003^{\circ}Z - 3.996$$

$$Z1 = -0.007^{\circ}X - 0.583^{\circ}Y - 0.021^{\circ}Z + 132.854$$

Original Joint Histogram



Final Joint Histogram



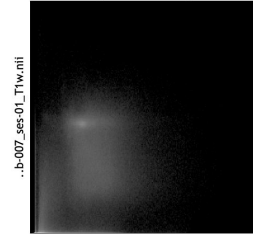
Normalised Mutual Information Coregistration

$$X1 = 0.003^{\circ}X - 0.021^{\circ}Y + 0.583^{\circ}Z + 21.946$$

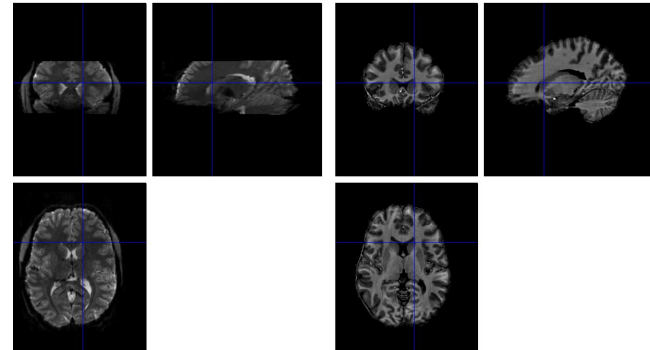
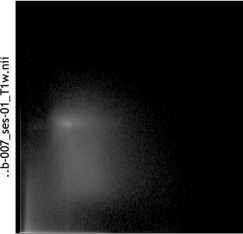
$$Y1 = 0.583^{\circ}X - 0.007^{\circ}Y - 0.003^{\circ}Z - 3.996$$

$$Z1 = -0.007^{\circ}X - 0.583^{\circ}Y - 0.021^{\circ}Z + 132.854$$

Original Joint Histogram



Final Joint Histogram



Two runs: Move T1 twice, or duplicate?

```
072_foodpicture_slicetiming.mat
072_realign_est&reslice_foodpicture.mat
072_realign_est&reslice_tasteimagine.mat
072_tatseimagine_slicetiming.mat
```

I moved on...

Current Module: Slice Timing

Help on: Slice Timing

Data	
. Session	194 files
. Session	194 files
. Session	194 files
. Session	194 files
Number of Slices	58
TR	2
TA	1.96551724137931
Slice order	1x58 double
Reference Slice	29
Filename Prefix	a

Current Item: Session

```
ta/ds004312/sub-072/ses-01/func/rsub-072_ses-01_task-foodpicture_run-04_bold.nii,1
ta/ds004312/sub-072/ses-01/func/rsub-072_ses-01_task-foodpicture_run-04_bold.nii,2
ta/ds004312/sub-072/ses-01/func/rsub-072_ses-01_task-foodpicture_run-04_bold.nii,3
ta/ds004312/sub-072/ses-01/func/rsub-072_ses-01_task-foodpicture_run-04_bold.nii,4
ta/ds004312/sub-072/ses-01/func/rsub-072_ses-01_task-foodpicture_run-04_bold.nii,5
ta/ds004312/sub-072/ses-01/func/rsub-072_ses-01_task-foodpicture_run-04_bold.nii,6
```

1st level analysis

Something's off....

Current Module: fMRI model specification

Help on: fMRI model specification

Directory /Users/annamarie/1stlevel/ds002995/sub-007/results

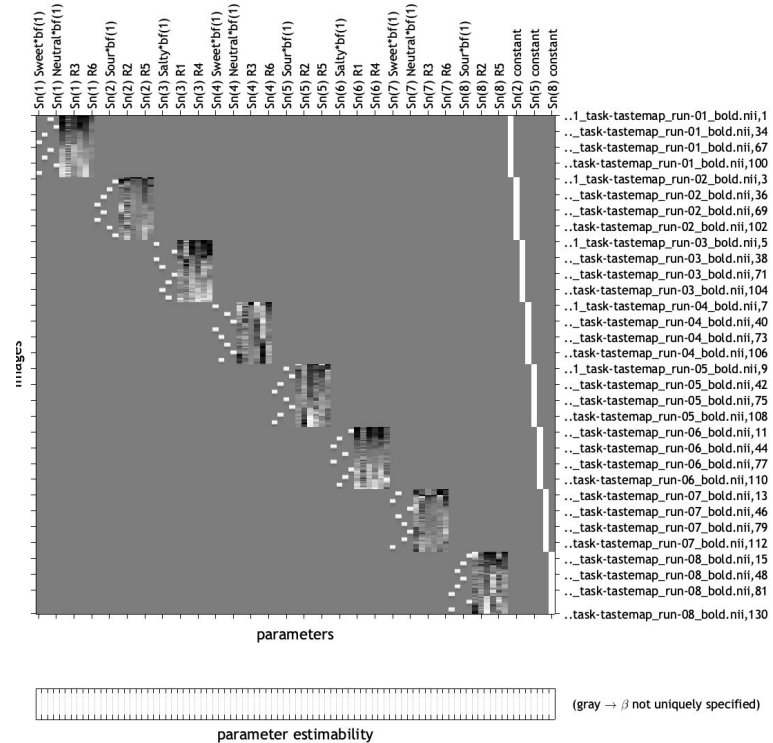
Timing parameters

. Units for design	Seconds
. Interscan interval	2.5
. Microtime resolution	68
. Microtime onset	7

Data & Design

. Subject/Session	
.. Scans	130 files
.. Conditions	
.. Multiple conditions	...arie/1stlevel/ds002995/sub-007/ses-01/func/Run1.mat
.. Regressors	
.. Multiple regressors	...unc/rp_sub-007_ses-01_task-tastemap_run-01_bold.txt
.. High-pass filter	128
. Subject/Session	
.. Scans	130 files
.. Conditions	
.. Multiple conditions	...arie/1stlevel/ds002995/sub-007/ses-01/func/Run2.mat
.. Regressors	
.. Multiple regressors	...unc/rp_sub-007_ses-01_task-tastemap_run-02_bold.txt

Statistical analysis: Design



Design description...

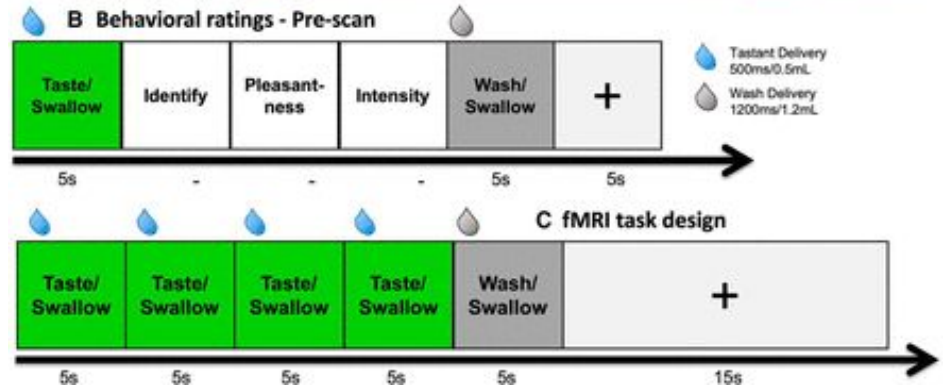
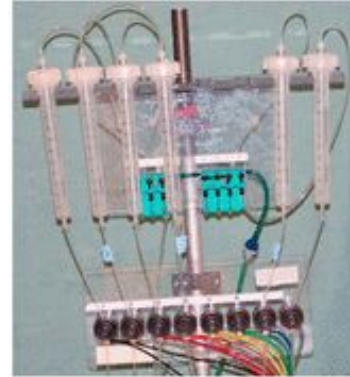
Basis functions : hrf
Number of sessions : 8
Trials per session : 4 4 4 4 4 4 4 4
Interscan interval : 2.50 [s]
High pass Filter : [min] Cutoff: 128 [s]
Global calculation : mean voxel value
Grand mean scaling : session specific
Global normalisation : None

1st study:

onset duration trial_type

5	20	sour
25	5	wash
45	20	neutral
65	5	wash
85	20	salty
105	5	wash
125	20	sweet
145	5	wash
165	20	sour
185	5	wash
205	20	salty
225	5	wash
245	20	neutral
265	5	wash
285	20	sweet
305	5	wash

*8 runs



1st level analysis

Checking .mat

```
>> load('Run1.mat')
```

```
>> whos
```

Name	Size	Bytes	Class	Attributes
ans	1x54	108	char	
durations	4x1	480	cell	
names	1x4	458	cell	
onsets	4x1	480	cell	

1st level analysis

Doing .mat again manually

```
>> names = {'sour', 'neutral', 'salty', 'sweet'};
```

```
onsets = {  
    [5, 165],      % sour  
    [45, 245],     % neutral  
    [85, 205],     % salty  
    [125, 285]     % sweet  
};
```

```
durations = {  
    [20, 20],      % sour  
    [20, 20],      % neutral  
    [20, 20],      % salty  
    [20, 20]       % sweet  
};|
```

```
>> save('Run1.mat', 'names', 'onsets', 'durations')
```

Nothing changed

Help on: fMRI model specification

Timing parameters

Units for design	Seconds
1	1
2	2
3	3
4	4
5	5
6	6
7	7
8	8
9	9
10	10
11	11
12	12
13	13
14	14
15	15
16	16
17	17
18	18
19	19
20	20
21	21
22	22
23	23
24	24
25	25
26	26
27	27
28	28
29	29
30	30
31	31
32	32
33	33
34	34
35	35
36	36
37	37
38	38
39	39
40	40
41	41
42	42
43	43
44	44
45	45
46	46
47	47
48	48
49	49
50	50
51	51
52	52
53	53
54	54
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56	56
57	57
58	58
59	59
60	60
61	61
62	62
63	63
64	64
65	65
66	66
67	67
68	68
69	69
70	70
71	71
72	72
73	73
74	74
75	75
76	76
77	77
78	78
79	79
80	80
81	81
82	82
83	83
84	84
85	85
86	86
87	87
88	88
89	89
90	90
91	91
92	92
93	93
94	94
95	95
96	96
97	97
98	98
99	99
100	100

. Interscan interval	2.5
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. Microtime resolution 68

. Microtime onset	7
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Data & Design

Subject/Session

```
.. Scans 130 files
```

.. Conditions

Multiple conditions ...arie/1stlevel/ds002995/sub-007/ses-01/func/Run1.ma

Regressors

```

-- Multiple regressors
...unc/rp sub-007 ses-01 task-tastemap run-01 bold.txt

```

High-pass filter	128
------------------	-----

Subject/Session	1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20	21	22	23	24	25	26	27	28	29	30	31	32	33	34	35	36	37	38	39	40	41	42	43	44	45	46	47	48	49	50	51	52	53	54	55	56	57	58	59	60	61	62	63	64	65	66	67	68	69	70	71	72	73	74	75	76	77	78	79	80	81	82	83	84	85	86	87	88	89	90	91	92	93	94	95	96	97	98	99	100
1	1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20	21	22	23	24	25	26	27	28	29	30	31	32	33	34	35	36	37	38	39	40	41	42	43	44	45	46	47	48	49	50	51	52	53	54	55	56	57	58	59	60	61	62	63	64	65	66	67	68	69	70	71	72	73	74	75	76	77	78	79	80	81	82	83	84	85	86	87	88	89	90	91	92	93	94	95	96	97	98	99	100

Scans	130 files
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11. Coating Conditions	100 times
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Multiple conditions	arie/1stlevel/ds002995/sub-007/ses-01/func/Bun2 ma
---------------------	--

Multiple conditions
Regressors

```
Multiple regressors      unc/rp_sub-007 ses-01 task-tasteman run-02 bold tv
```

(gray $\rightarrow \beta$ not uniquely specified)

parameter estimability

Design description...

Basis functions : hrf
 Number of sessions : 8
 Trials per session : 4 4 4 4 4 4 4 4
 Interscan interval : 2.50 {s}
 High pass Filter : [min] Cutoff: 128 {s}
 Global calculation : mean voxel value
 Grand mean scaling : session specific
 Global normalisation : None

Future Direction

- Manually input conditions
- Checks if smoothing/additional steps needed before model specification
- 1st level > 2nd level analysis
- Move beyond visual inspection.
- Make a T shirt with sweet contrast

