Neural reuse in the evolution and development of the brain

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Where we are

20+ years of efforts in non-invasive functional brain imaging

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Tens of thousands of studies =

A huge scientific opportunity to revisit some of the "big" questions

Overall functional architecture













Notice what's happening:

- New functions develop by incorporating parts used in existing functions.
- Over time, many local circuits get reused in different cognitive functions.
- That reuse may be in different cognitive domains

This is a "component reuse" or "*buildingblock*" model of brain evolution.

Components evolved for one cognitive purpose are "exapted" for later uses.

However, the original functionality is not lost (reuse not driven by plasticity)

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Overall functional architecture



Neural reuse means that:

I. Local neural circuits are typically redeployed in support of multiple cognitive functions, across traditional task-domain boundaries

II. Redeployed circuits will have the same "working" (but a different "use") in each of the functional complexes they support

III. Task domains will differentially activate not *regions*, but *networks*

<u>Interactions between</u> regions will be a better indication of function than <u>actions of</u> regions

IV. More recent cognitive functions will utilize more, and more widely scattered local circuits

Neural reuse

- I. Local circuits are typically redeployed in support of multiple cognitive functions across traditional task-domain boundaries
- II. Redeployed circuits have the same "working" in each of the functional complexes they support
- III. Task domains will be differentiated at the network level
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Implication I: Selectivity

The primary guiding idealization of the functional organization of the brain for at least 50 years.

Brain regions selectively respond to

- Specific input classes
- Tasks and task categories

Functional Selectivity



Penfield's Homunculus, 1954

Domain Selectivity



Brodmann map showing functional domains

NICAM Database

NICAM database contains:

- 2,603 experiments from 824 journal articles investigating over 70 different task domains
- **21,553** regional activations
- Only: Whole-brain analysis, postsubtraction activations

Empirical Review of Imaging Experiments

Investigated 11 task domains:

- three action domains (execution, inhibition and observation)
- two perceptual domains (vision and audition)
- six "cognitive" domains (attention, emotion, language, working memory, semantic memory and reasoning).

Empirical Review of Imaging Experiments

Gives 1,469 experiments collectively reporting 10,701 eligible activations.

Procedure:

- 1. divide the brain into regions
- 2. assign each activation to a region.
- 3. measure the diversity of activations

Diversity measure

Borrowed from the Census bureau

$$DV = \sqrt{\frac{\sum_{i=1}^{k} (Cat_i - mean)^2}{k}}$$

Results with 66 cortical + 12 sub-cortical

Overall average diversity: 0.70 (sd 0.12)

Overall cortical diversity: 0.71 (sd 0.11)

Overall sub-cortical diversity 0.63 (sd 0.17)

Regions active in an average of **95** tasks across **9** different domains

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Results with 66 cortical + 12 sub-cortical

Putting that in terms of ratio of posterior to prior probability (Poldrack 2006):

Average Bayes factor = 1.26

Even the *best* evidence is fairly weak: average MAX Bayes factor = 3.14 (and this presumes you know what the best evidence is)

Left inferior temporal cortex has a diversity of 0.70

Domain	Proportion of activations
Action execution	0.06
Action imagination	0.06
Action inhibition	0.00
Visual perception	0.07
Audition	0.02
Attention	0.02
Semantic memory	0.12
Working memory	0.12
Emotion	0.06
Reasoning	0.12
Language	0.34





0.89 - 0.84	
0.84 - 0.80	
0.80 - 0.75	
0.75 - 0.70	
0.70 - 0.66	
0.66 - 0.61	
0.61 - 0.56	
0.56 - 0.52	
0.52 - 0.47	
0.47 - 0.42	
0.42 - 0.37	
0.37 - 0.33	
0.33 - 0.28	
0.28 - 0.23	
0.23 - 0.19	
0.19 - 0.14	
0.14 - 0.09	
0.09 - 0.05	
0.05 - 0.00	









0.89 - 0.84
0.84 - 0.80
0.80 - 0.75
0.75 - 0.70
0.70 - 0.66
0.66 - 0.61
0.61 - 0.56
0.56 - 0.52
0.52 - 0.47
0.47 - 0.42
0.42 - 0.37
0.37 - 0.33
0.33 - 0.28
0.28 - 0.23
0.23 - 0.19
0.19 - 0.14
0.14 - 0.09
0.09 - 0.05
0.05 - 0.00







0.89-0.87
0.87-0.85
0.85-0.83
0.83-0.81
0.81-0.79
0.79-0.77
0.77 – 0.75
0.75-0.73
0.73-0.71
0.71-0.69
0.69-0.67
0.67 - 0.65
0.65-0.63
0.63-0.61
0.61-0.59
0.59-0.57
0.57-0.55
0.55-0.53
< 0.53









0.89-0.87
0.87-0.85
0.85-0.83
0.83-0.81
0.81-0.79
0.79-0.77
0.77-0.75
0.75-0.73
0.73-0.71
0.71-0.69
0.69-0.67
0.67-0.65
0.65-0.63
0.63-0.61
0.61-0.59
0.59-0.57
0.57-0.55
0.55-0.53
< 0.53





With 998 cortical + 54 sub-cortical Results for 574 + 21 with > 5 activations

Overall average diversity: 0.52 (sd 0.13)

Overall cortical diversity: 0.52 (sd 0.13)

Overall sub-cortical diversity 0.59 (sd 0.12)

Regions active in an average **10** tasks over **4** different domains

Results with 998 cortical + 54 sub-cortical

Putting that in terms of ratio of posterior to prior probability (Poldrack 2006):

Average Bayes factor = 2.72

Even the **best** evidence is only moderately strong: average MAX Bayes factor = 5.54 (and this presumes you know what the best evidence is)

Sub-region of right supramarginal gyrus with diversity of 0.52

Domain	Proportion of activations
Action execution	0.27
Action imagination	0.42
Action inhibition	0.00
Visual perception	0.00
Audition	0.00
Attention	0.00
Semantic memory	0.00
Working memory	0.11
Emotion	0.00
Reasoning	0.20
Language	0.00

Selectivity

Provisional conclusion I:

Local circuits typically support multiple cognitive functions across traditional task-domain boundaries

Neural reuse



Local circuits are typically redeployed in support of multiple cognitive functions across traditional task-domain boundaries

- II. Redeployed circuits have the same "working" in each of the functional complexes they support
- III. Task domains will be differentiated at the network level
- IV. More recent cognitive functions will utilize more, and more widely scattered local circuits

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Overall functional architecture




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III: Networks or regions?

- "Functional connectivity" analysis of the same data set.
- Only 10 domains insufficient data for functional connectivity study of action inhibition

Functional connectivity

- 1) Calculate chance probability (**Q**) of coactivation for each BA pair
- In each domain, determine observed probability (*K*) of co-activation of each BA pair
- Where there is a significant difference between *Q* and *K* (X²), this is considered a "functional connection"

Functional architecture

The results of functional connectivity analyses can be represented as graphs

What's a graph?

Graphs and graph theory

Graph: a set of "nodes" joined by "edges"

Examples: Airline route map Social network Website map Brain adjacency map













Attention

Comparing Domain Complexes

Can compare many things, for instance:

- -Node overlap
 - Indicates local circuits shared by different domain complexes
- -Edge overlap
 - Indicates functional connectivity/ cooperation shared by different domain complexes

Comparing Domain Complexes

-Network topology

• May give clues about nature of function implementation

-Degree

 Indication of how many other regions a brain region typically cooperates with







Attention

Overall functional architecture



Node vs. Edge Overlap

Use Dice's coefficient: $2(o_{1,2})/(n_1+n_2)$

Predictions:

- Modularity: e, n
- Holism: E, N
- Reuse: e, N



55







p << 0.001Workshop on Developmental Homology

But . . .

Maybe this result is just an artifact

- Given a small number of nodes (111)
- Large number of possible edges (6105)
- Get high node overlap and low edge overlap just by chance



p << 0.001



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Some implications

No use selectivity means NO REVERSE INFERENCE, PLEASE

Some implications

Regions of the brain should have differential, but measurable *burden*.

This is perhaps correlated with diversity?

Some implications

Assigning computational/cognitive "workings" to local circuits will require cross-domain modeling

But is it *Homology*?

Dagher et al. 1999: the reuse of motor circuits for

- action sequence planning
- abstract planning

This could be a serial homology, with shared developmental precursor, and "special qualities"

But is it *Homology*?

Although, note there is no need for any "copying" of the shared structure

The selection pressure can operate on the *networks*, which differ

But is it *Homology*?

On the other hand,

Since what differentiates the two is precisely the *different connections* of the shared structure,

Is this a *bone*, or a *turtle shell*?

Finger gnosia and math

•*Finger gnosia* is the ability to distinguish, in the absence of visual feedback, which fingers have been lightly touched.

•Developmentally, finger gnosia has been found to predict mathematical ability

	Digit Recognition	Next Number	Place Value	Numeration	Calculation
Finger Gnosia	.19*	.22**	.18*	.28**	.19*


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But is it *Homology*?

In this case the shared working appears to be a particular sort of storage resource: an array of pointers

Here any inheritance of functional character is much less clear; it is simply reuse of an available storage system

But is it *Homology*?

And, of course, here again, differential function is achieved precisely in terms of *differential connections* to other "organs"

But is it *Homology*?

On the other hand,

It does appear that the systems *fail* in similar ways

Does that count as a special quality?

So is it *Homology*?



(I don't have a conclusion)

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