

Linking retinotopic fMRI mapping and anatomical probability maps of human occipital areas V1 and V2



Ritzl A^{1,2}, Specht K^{1,3}, Lie C-H¹, Mohlberg H¹, Wohlschläger A⁴, Bente K¹, Pietrzyk U¹, Stöcker T¹, Zilles K^{1,5}, Amunts K¹, Fink GR^{1,3}

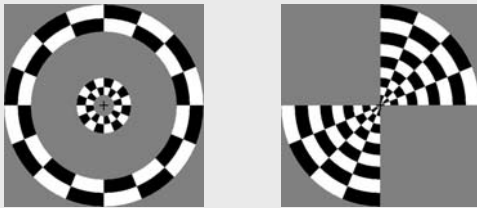
¹Institut für Medizin, Forschungszentrum Jülich, Germany; ²Neurologie, Nuklearmedizin und Radiologie, Technische Universität München, Germany;

³Neurologische Klinik, Universitätsklinikum der RWTH Aachen, Germany; ⁴Arbeitsbereich Psychologie, Max Planck Institut für Kognitions- und Neurowissenschaften, München, Germany; ⁵C. & O. Vogt - Hirnforschungsinstitut, Heinrich-Heine-Universität Düsseldorf, Germany

Introduction It has been known for long that the cytoarchitectonic organization of the human visual cortex is associated with functional organization. **We here compare retinotopic fMRI mapping and anatomical probability maps** derived from post-mortem studies of cytoarchitecture to establish whether or not the two methods lead to converging results. To allow for a comparison, 3D-probability maps for occipital areas V1 and V2 were created from functional data and compared to cytoarchitectonically defined probability maps of V1 and V2.

Methods

- ❖ 12 subjects, fMRI and T1-weighted measurement on Siemens Sonata 1.5T-scanner
- ❖ standard periodic retinotopic stimuli [1]:



- ❖ preprocessing in spm99 including normalization to ICBM 152 brain
- ❖ calculation of **3D-fieldmaps** according to [2]
- ❖ creation of probability maps of areas with positive (V1) and negative (V2) field sign as mean images
- ❖ **comparison to anatomical probability maps of BA17 and BA18** from cytoarchitectonic analyses of post-mortem brains [3]
- ❖ **statistical test** using a-priori measure κ [4]:

$$\kappa = 1 - NE(H1)/NE(H0),$$
 with $NE(H1)$ = number of errors under $H1$ and $NE(H0)$ = number of errors under $H0$, where $H1$ = ‘a voxel is only in V1 if it is also in BA17’ and $H0$ = ‘a voxel being in V1 or BA17 is independent’ or equivalently for V2 and BA18

($\kappa = 1$ means that a brain region can only be in the functionally defined retinotopic area if it is also in the anatomically defined one. $\kappa = 0$ means that being in the functionally defined area and the anatomically defined could equally be independent)

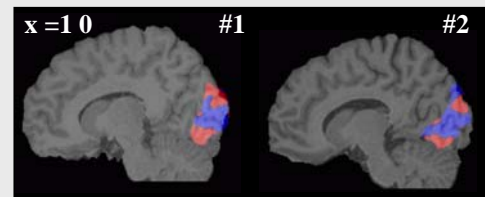


Fig.1: examples of field sign maps (blue = +1, red = -1)

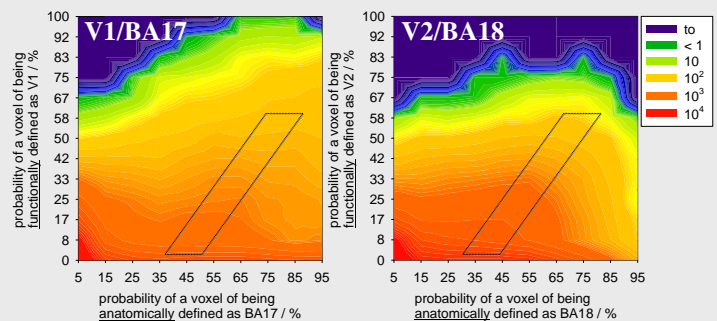


Fig.2: correlation plot of functional and anatomical data (voxel numbers on logarithmic scale)

Results V1: maximum probability found functionally was 92 % in an (overlap-)volume of 104 mm³ with centres of gravity in the right hemisphere (RH) at 10, -80, 0 and in the left hemisphere (LH) at -8, -82, -1.

Corresponding anatomical map BA17: the respective coordinates were 15, -82, -1 (RH) and -10, -84, -2 (LH).

V1 and BA17: $\kappa = 0.99$.

V2: peak probability of 67 % in 196 mm³ with centres of gravity at 11, -95, 15 (RH) and -13, -97, 10 (LH)

Corresponding anatomical map BA18: 16, -85, 1 (RH), and -10, -89, 0 (LH)

V2 and BA18: $\kappa = 0.76$.

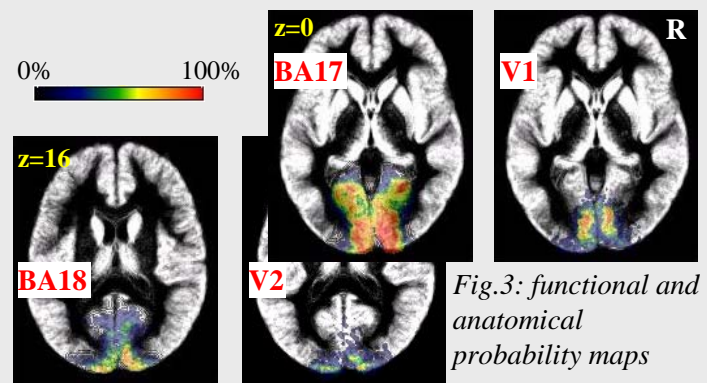


Fig.3: functional and anatomical probability maps

Discussion Comparison of functionally derived probability maps of V1 and V2 with cytoarchitectonically derived anatomical probability maps shows a **good spatial agreement**. Observed discrepancies are likely to reflect restrictions in the maximum angle of visual stimulation during the functional measurements and inter-subject variability.

[1] Engel et al. (1997) Cereb. Cortex 7:181-192.

[2] Dumoulin et al. (2003) NeuroImage 18:576-587.

[3] Amunts et al. (2000) NeuroImage 11:66-84.

[4] Cohen J (1960) Educational and Psychological measurement 20:37-46.