## Data from: Inhibition decorrelates visual feature representations in the inner retina

This dataset contains raw and processed data acquired using 2-photon imaging with iGluSnFr of mouse bipolar cells using a standardized set of stimuli as described in Franke et al. (2017) after application of various drugs. Visualization functions for use with MATLAB are available from www.retinal-

Variable name	Dim 1	Dim 2	Dim 3	Dim 4	Remarks	
gchirp_drug_avg gchirp_drug_qi	#samples #rois	#rois			Median of all repeats, scaled to max(abs(x))=1  Quality index as defined in paper	global chirp
lchirp_drug_avg lchirp_drug_qi	#samples #rois	#rois			Median of all repeats, scaled to max(abs(x))=1 Quality index as defined in paper	local chirp
step_drug_avg step_drug_qi	#samples #rois	#rois			Median of all repeats, scaled to max(abs(x))=1 Quality index as defined in paper	
rf_drug_tc rf_drug_map rf_drug_gauss_mean			15 #rois 2 2		Time kernel of RF Spatial map of RF 2D mean of gauss fit to the spatial map	
<pre>rf_drug_gauss_std rf_drug_size</pre>	#rois #rois		2		2D std of gauss fit to the spatial map 2-std-ellipse area as receptive field size	
cs_drug_rf	#samples		10 #rois		Center-surround maps using the annulus stimulus Center kernel (not identical to those used in the paper - the analysis in the	
cs_drug_center	#samples	#rois			paper was implemented in IGOR and is available upon request)	
cs_drug_surround	#samples	#rois			Surround kernel (not identical to those used in the paper - the analysis in the paper was implemented in IGOR and is available upon request)	9
drug	#rois				1: Gabazine, 2: Tpmpa, 3: Strychnine, 4: L-AP4, NaN: no drug condition for	this ROI

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