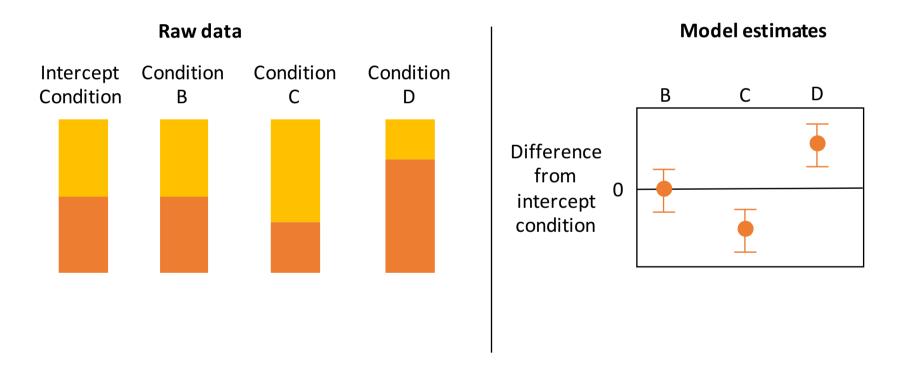
This document was written to explain how the munltinomial model works in general, and was produced before the data was finalised. Readers should refer to the paper for the final numerical results.

Interpretation of multinomial models

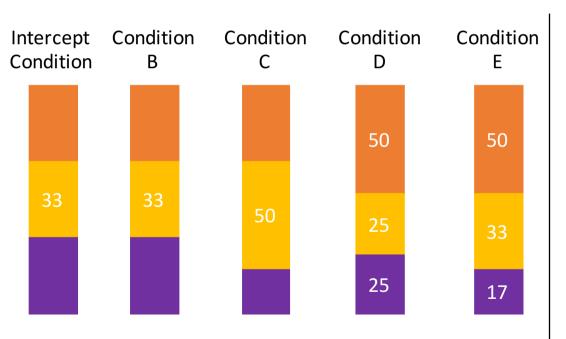
If a variable has two levels, we just need to estimate one number for each condition: how much one of the levels changes.

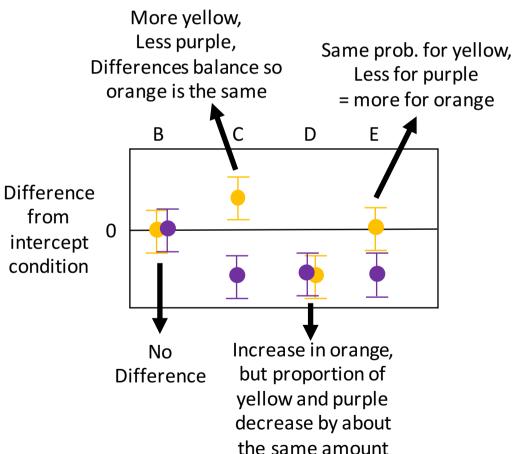
Here's an imaginary example:



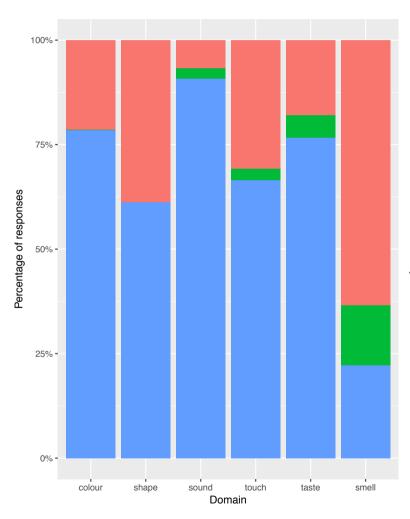
Interpretation of multinomial models

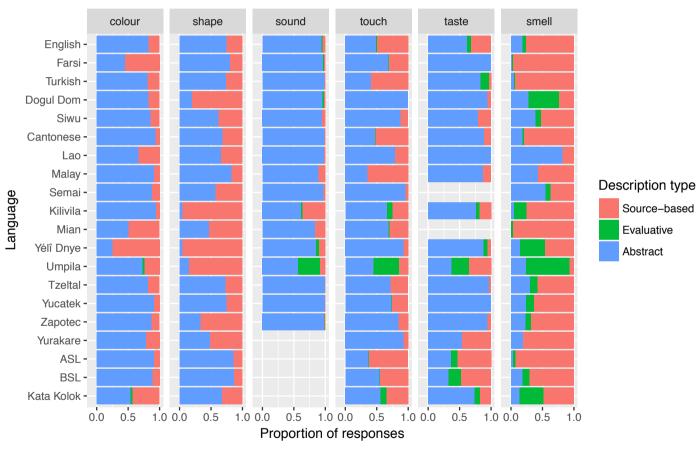
Our variable has 3 levels. One level and one condition is chosen as the intercept. We need to estimate two numbers: change in intercept level and change in one other level (the final level is implied logically from the other two)





Raw proportions





Results

Intercept is abstract terms for colours.

Output of the	
MCMC model	

	<pre>post.mean</pre>	1-95% CI	u-95% CI	eff.samp	pMCMC
traitSAE.E	-6.73	-7.480	-5.95	1160.00	0.00002
traitSAE.S	-1.87	-2.580	-1.16	47400.00	0.00004
domainshape	-7.99	-12.500	-1.96	2.76	0.00280
domainsound	1.79	0.691	2.91	2830.00	0.00228
domaintouch	2.70	1.430	3.99	4410.00	0.00028
domaintaste	2.80	1.310	4.35	4730.00	0.00052
domainsmell	6.70	5.590	7.78	2560.00	0.00002
traitSAE.S:domainshape	9.53	3.920	13.70	2.50	0.00002
traitSAE.S:domainsound	-3.81	-4.190	-3.43	371.00	0.00002
traitSAE.S:domaintouch	-2.23	-2.630	-1.84	381.00	0.00002
traitSAE.S:domaintaste	-3.53	-3.990	-3.09	414.00	0.00002
<pre>traitSAE.S:domainsmell</pre>	-3.15	-3.480	-2.83	126.00	0.00002

We can plot these estimates and confidence intervals \dots

Results

Estimate (mean of distribution taken from the posterior)

Measure of the independence of the samples from the MCMC

Probability that the estimate is not different from zero

Intercept is abstract terms for colours.

Confidence intervals

E and S are less likely than abstract in general

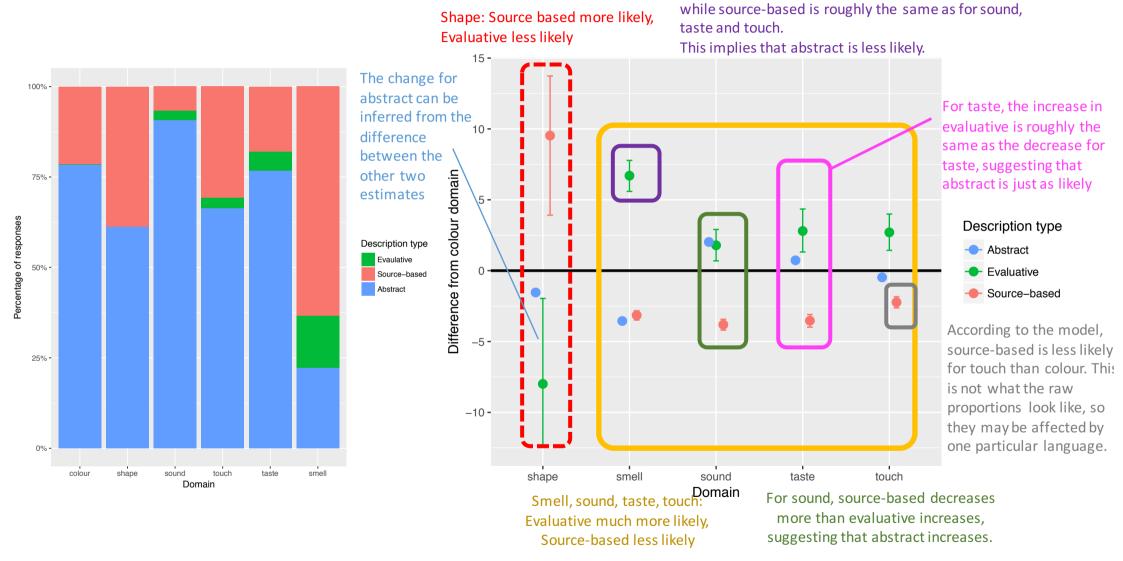
Estimates for Evaluative descriptions by domain

Estimates for Sourcebased descriptions by domain

				•	
	post.mean	1-95% CI	u-95% CI	eff.samp	pMCMC
traitSAE.E	-6.73	-7.480	-5.95	1160.00 0	0.00002
traitSAE.S	-1.87	-2.580	-1.16	47400.00 6	0.00004
domainshape	-7.99	-12.500	-1.96	2.76 6	0.00280
domainsound	1.79	0.691	2.91	2830.00 6	0.00228
domaintouch	2.70	1.430	3.99	4410.00 6	0.00028
domaintaste	2.80	1.310	4.35	4730.00 6	0.00052
domainsmell	6.70	5.590	7.78	2560.00 0	0.00002
traitSAE.S:domainshape	9.53	3.920	13.70	2.50 6	0.00002
traitSAE.S:domainsound	-3.81	-4.190	-3.43	371.00 6	0.00002
traitSAE.S:domaintouch	-2.23	-2.630	-1.84	381.00 6	0.00002
traitSAE.S:domaintaste	-3.53	-3.990	-3.09	414.00 6	0.00002
traitSAE.S:domainsmell	-3.15	-3.480	-2.83	126.00 6	0.00002

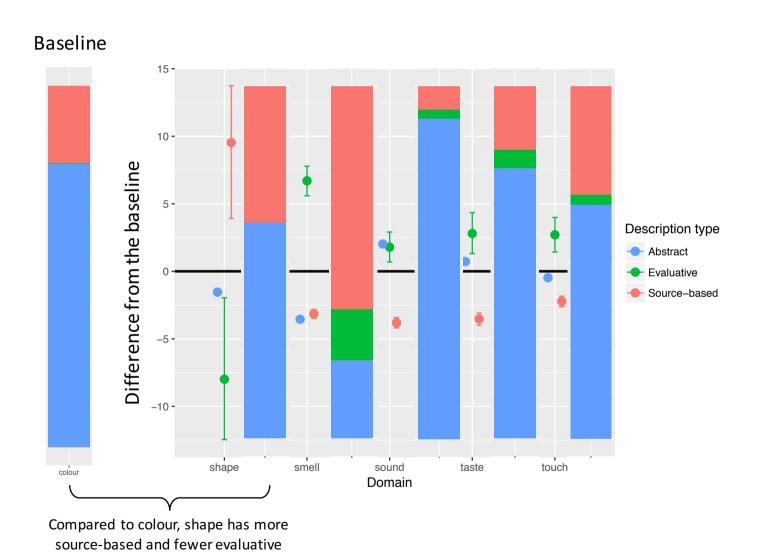
We can plot these estimates and confidence intervals ...

Intercept is abstract terms for colours.



For smell: source-based evaluative is much higher,

Intercept is abstract terms for colours.



Note: This run done before clean-up of shape typos

Results (intercept is source-based)

Abstract terms more likely than source based, Evaluative less likely than source based

Abstract responses for taste not different to abstract responses for colour

urce baseu	post.mean	1-95% CI	u-95% CI	eff.samp	pMCMC
traitSAE.A	1.730	1.330	2.140	88900.0	0.00001
traitSAE.E	-4.580	-5.090	-4.060	641.0	0.00001
domainsound	2.020	1.620	2.430	18900.0	0.00001
domaintaste	0.258	-0.435	0.961	58100.0	0.46500
domaintouch	-0.718	-1.210	-0.209	75100.0	0.00470
domainshape	-1.000	-1.370	-0.633	66300.0	0.00001
domainsmell	-3.050	-3.510	-2.590	69700.0	0.00001
traitSAE.E:domainsound	2.300	1.910	2.670	457.0	0.00001
<pre>traitSAE.E:domaintaste</pre>	3.280	2.860	3.680	746.0	0.00001
traitSAE.E:domaintouch	2.780	2.390	3.160	610.0	0.00001
traitSAE.E:domainshape	-4.050	-7.230	-1.260	14.9	0.00001
<pre>traitSAE.E:domainsmell</pre>	5.840	5.480	6.170	303.0	0.00001

