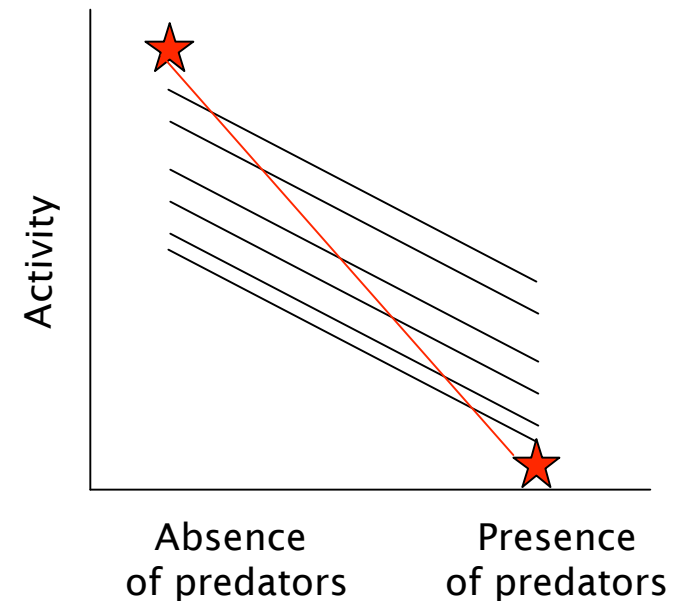
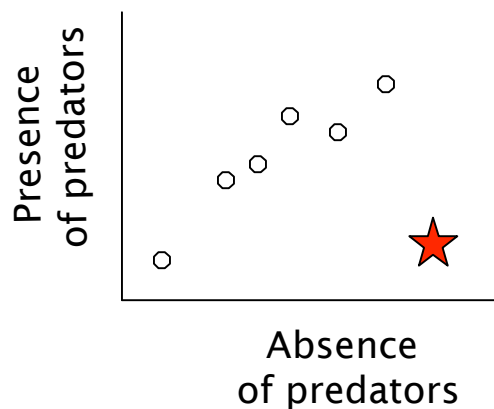


Personality is mysterious from an evolutionary point of view, for three reasons

1. Why should individuals behave consistently, either through time or across functional contexts?



Personality suggests cleverness?

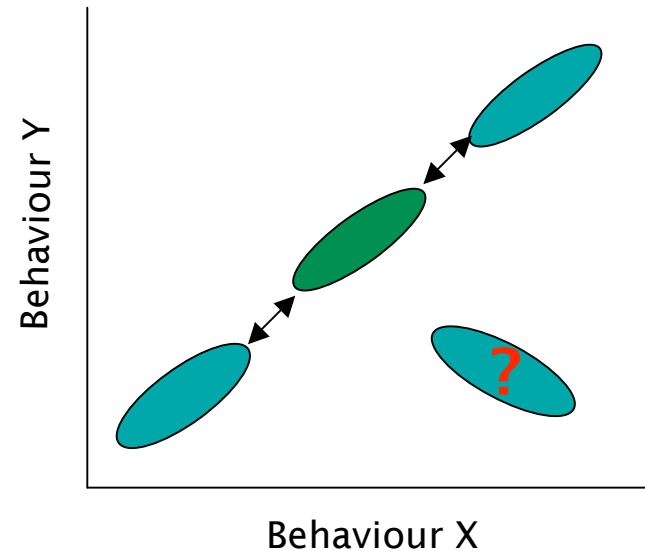
Personality is mysterious from an evolutionary point of view, for three reasons

1. Why should individuals behave consistently, either through time or across functional contexts?
2. Why should individuals *differ* in how they behave?

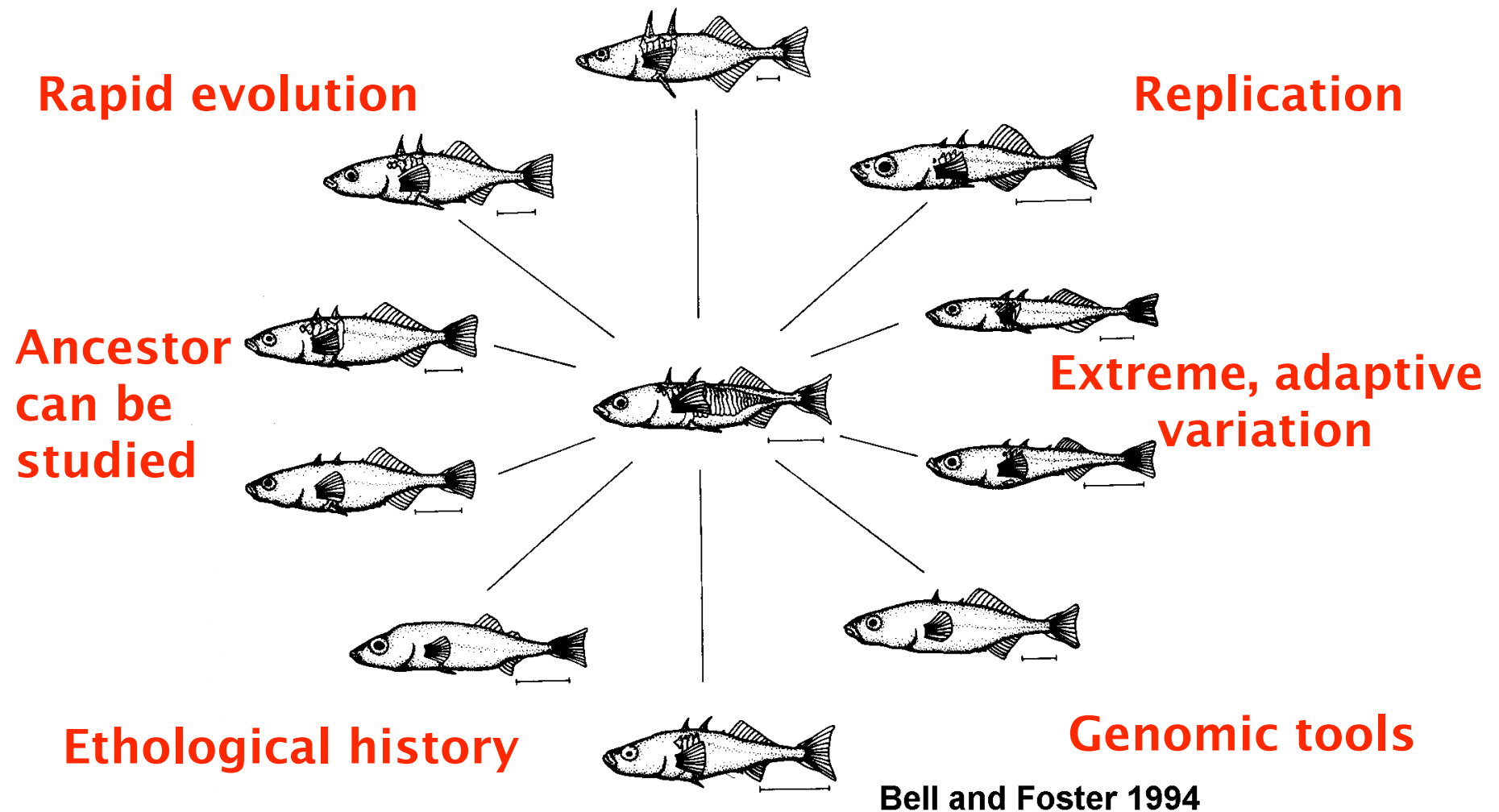
Natural selection erodes heritable variation that is related to fitness

Personality is mysterious from an evolutionary point of view, for three reasons

1. Why should individuals behave consistently either through time or across functional contexts?
2. Why should individuals *differ* in how they behave?
3. Evolutionary constraints

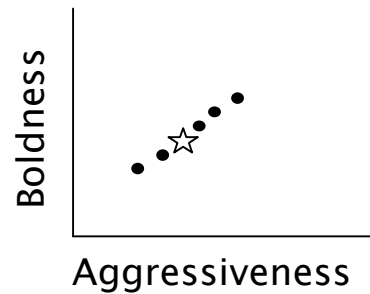


What can the radiation of threespined sticklebacks tell us about the evolution of personality?

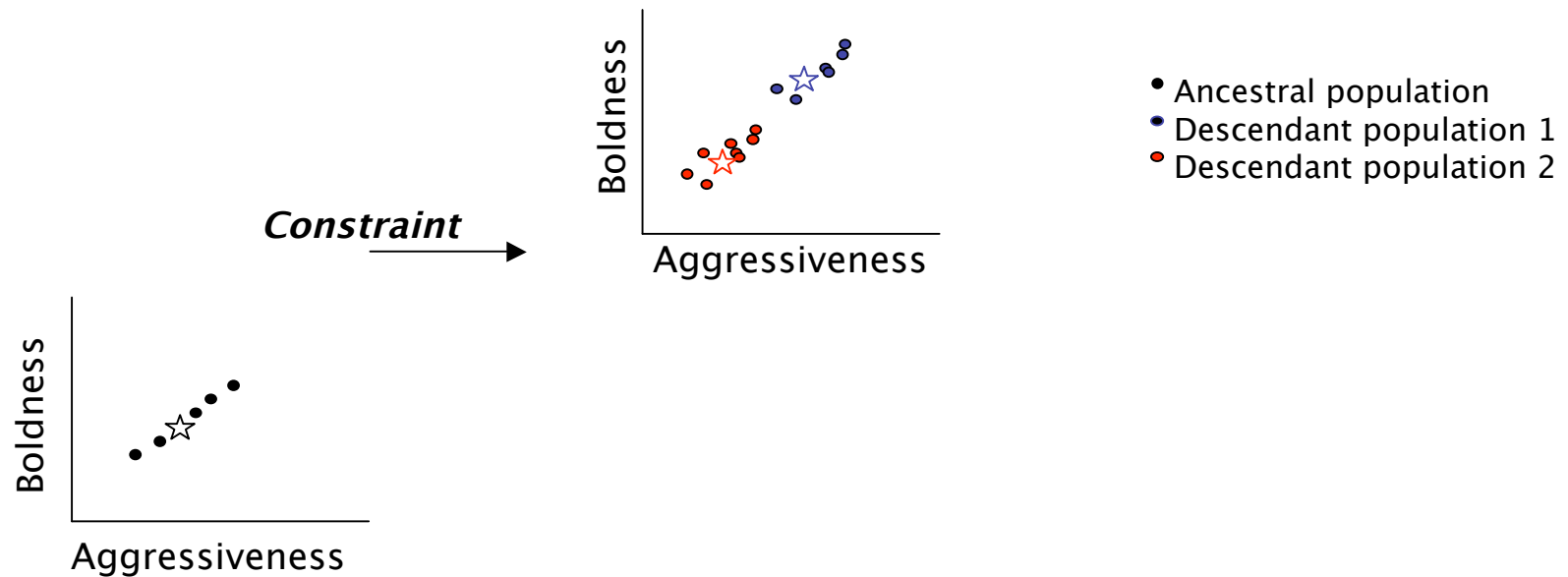


The evolutionary fate of correlated traits

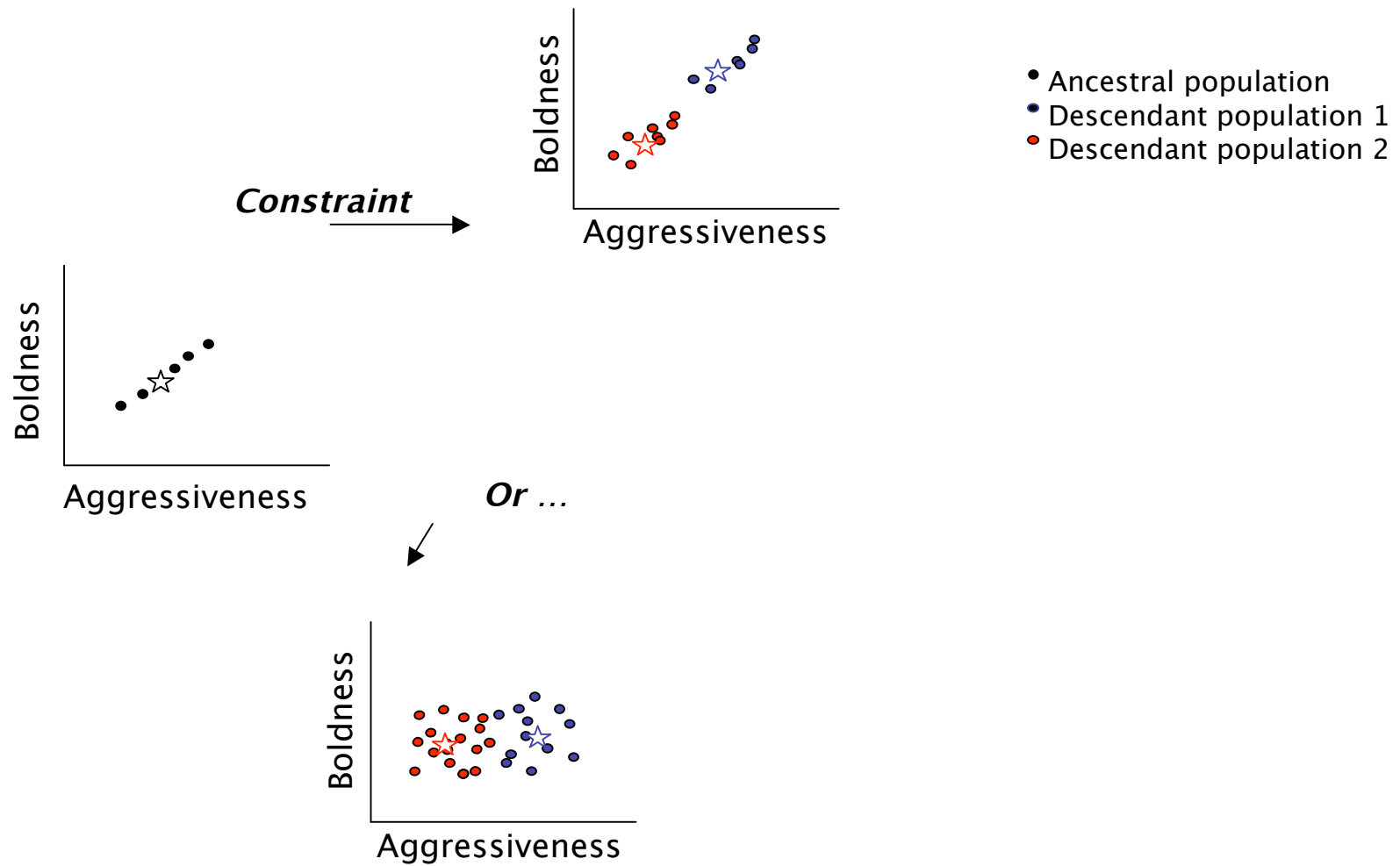
- Ancestral population
- Descendant population 1
- Descendant population 2



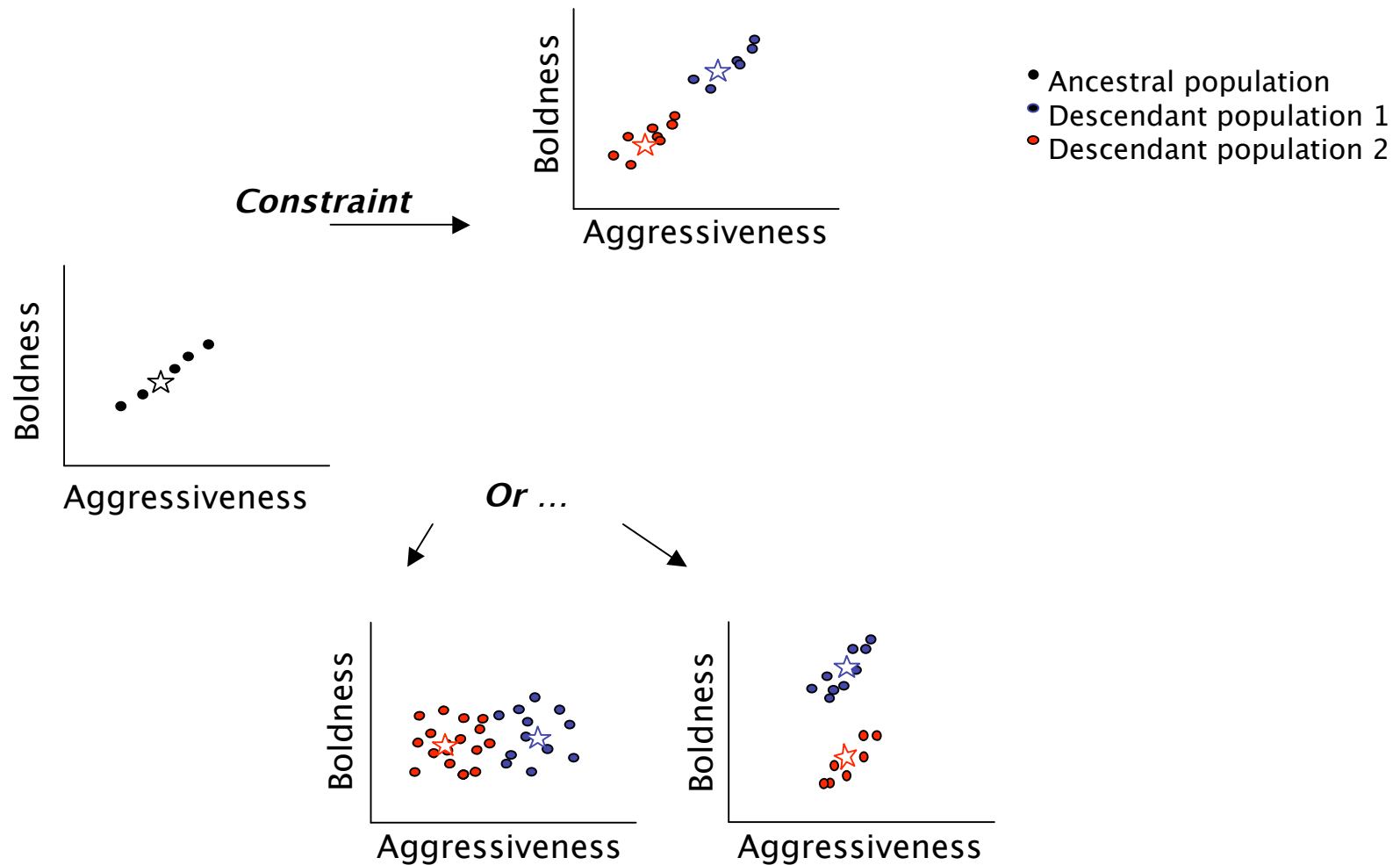
The evolutionary fate of correlated traits



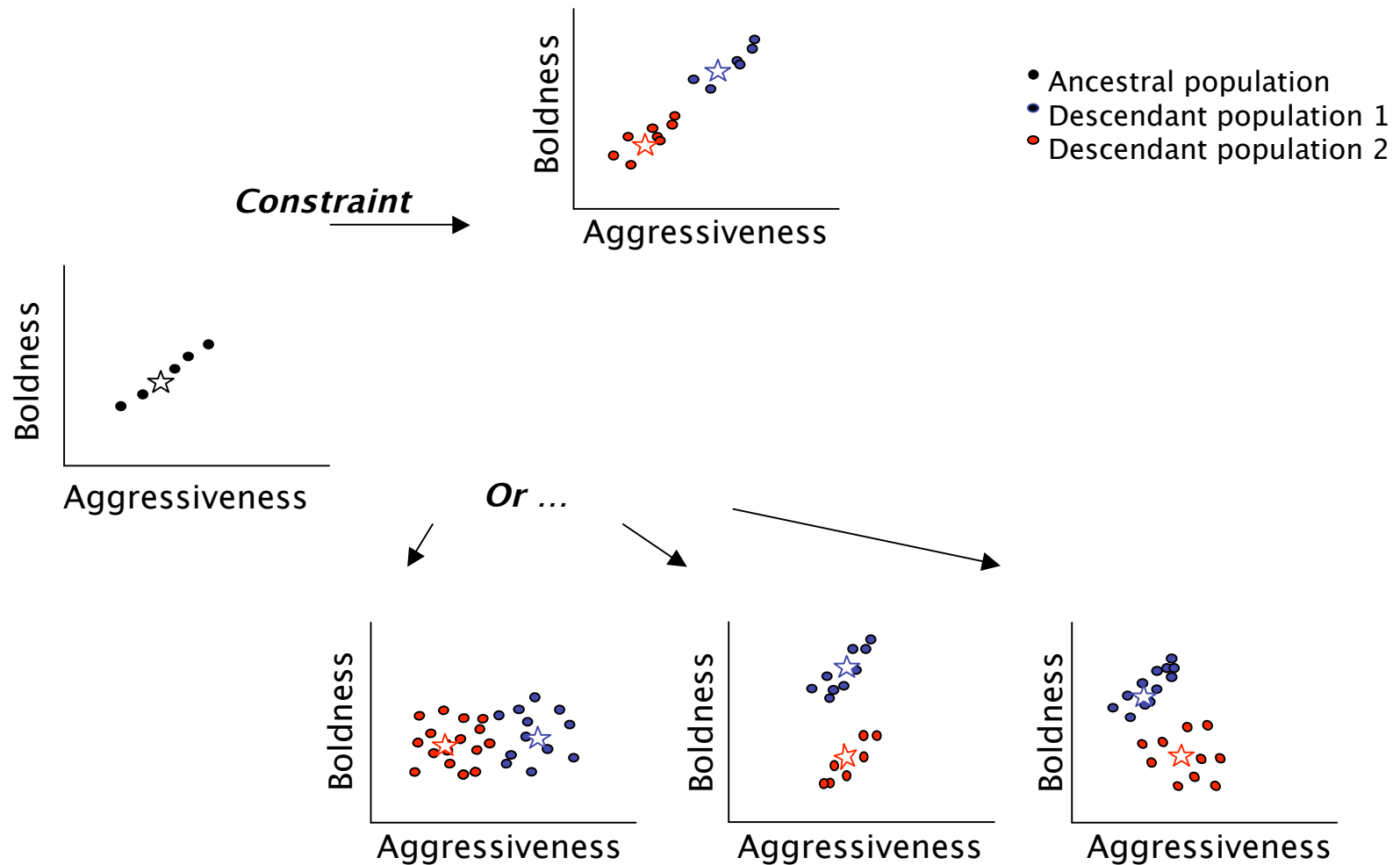
The evolutionary fate of correlated traits



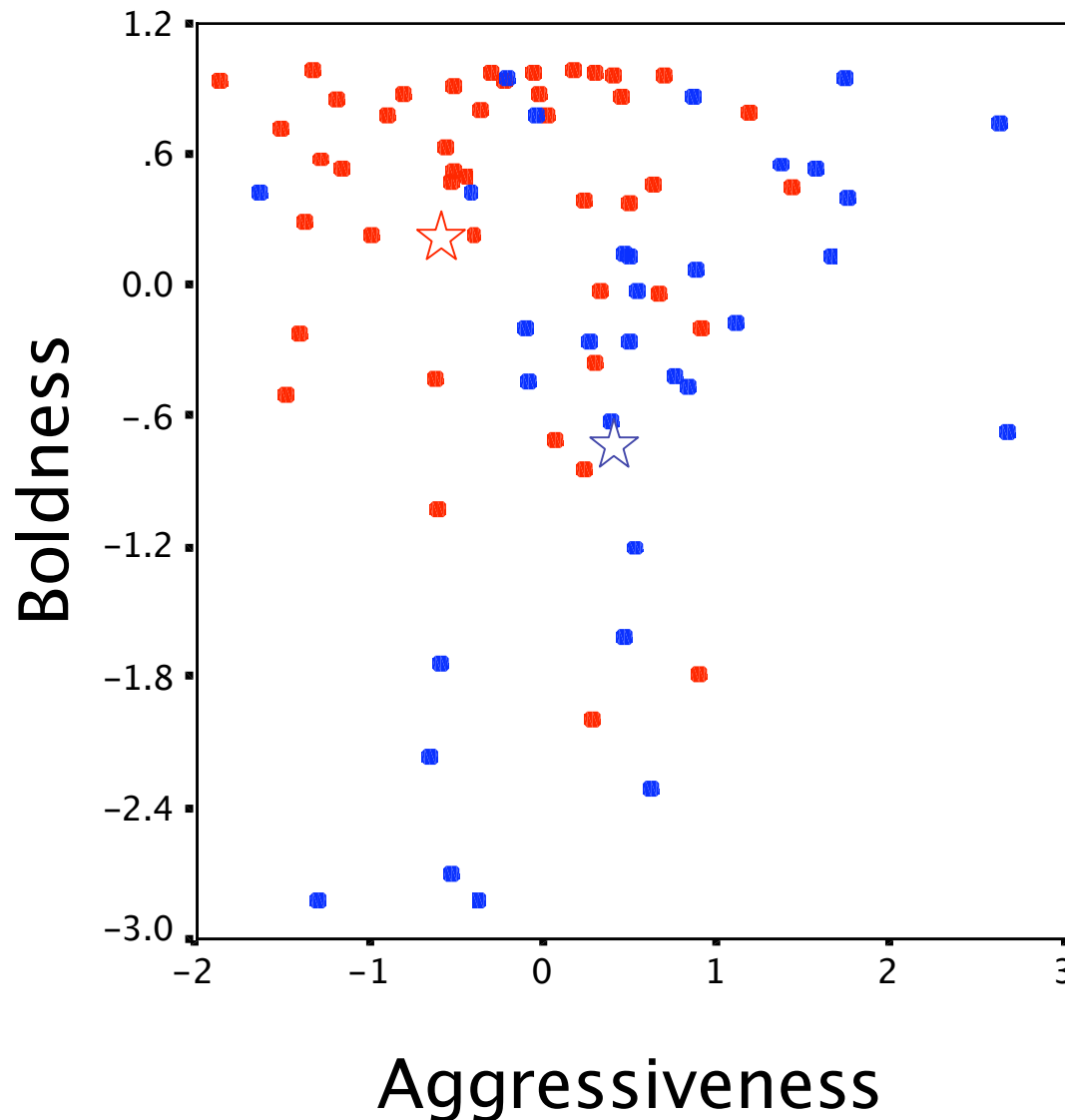
The evolutionary fate of correlated traits



The evolutionary fate of correlated traits



Boldness and aggressiveness were genetically correlated in one population but not another (Bell 2005 *JEB*)



ALSO: Boldness and aggressiveness were uncoupled over ontogeny
(Bell & Stamps 2004 *An Behav*)

● Navarro: $r=0.52$, $n=29$, $P<0.01$
● Putah: $r=0.13$, $n=42$, *NS*

Can experimental exposure to real predation risk generate the syndrome?

With Andy Sih, UC Davis



Can experimental exposure to real predation risk generate the syndrome?

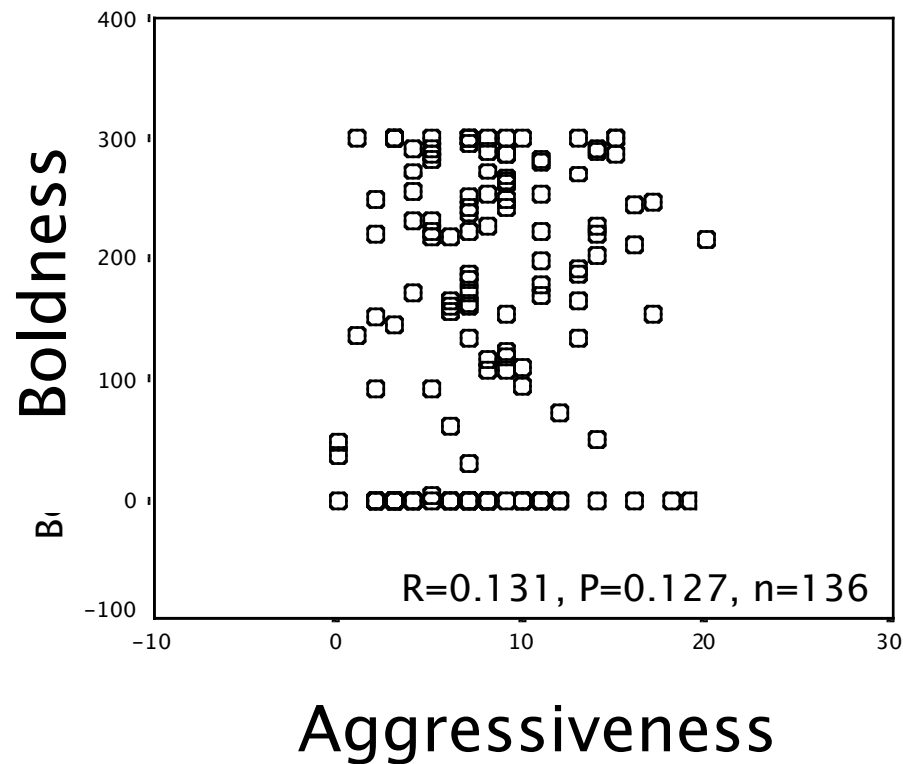
With Andy Sih, UC Davis



- Measured boldness (time spent eating under predation risk) and aggressiveness (orienting and attacking a conspecific) on individually-marked sticklebacks.
- Placed groups (n=34/pool) of individually-marked fish in seminatural pools (n=4 replicate pools) with trout. Control: No trout.
- Removed trout when 1/2 of the sticklebacks had been consumed.
- Re-measured survivors.

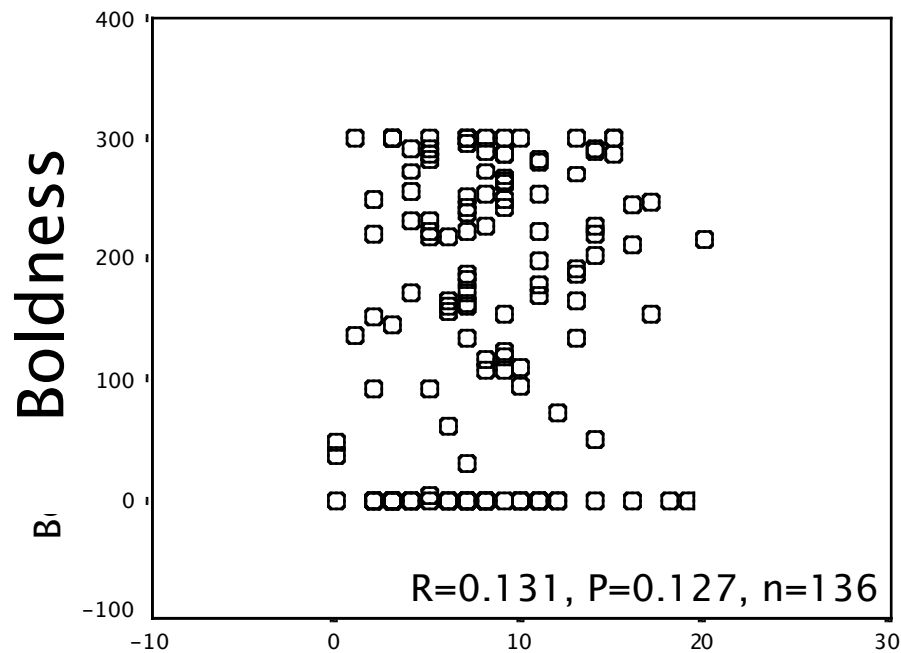
Boldness and aggressiveness were not correlated with each other 'before'

BEFORE



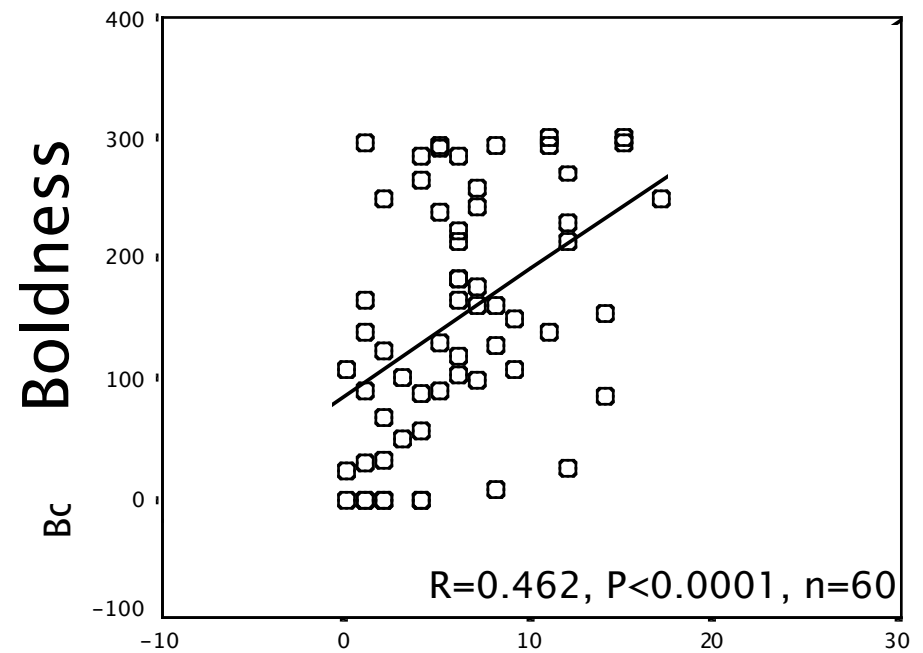
Predation generated the syndrome!

BEFORE



Aggressiveness

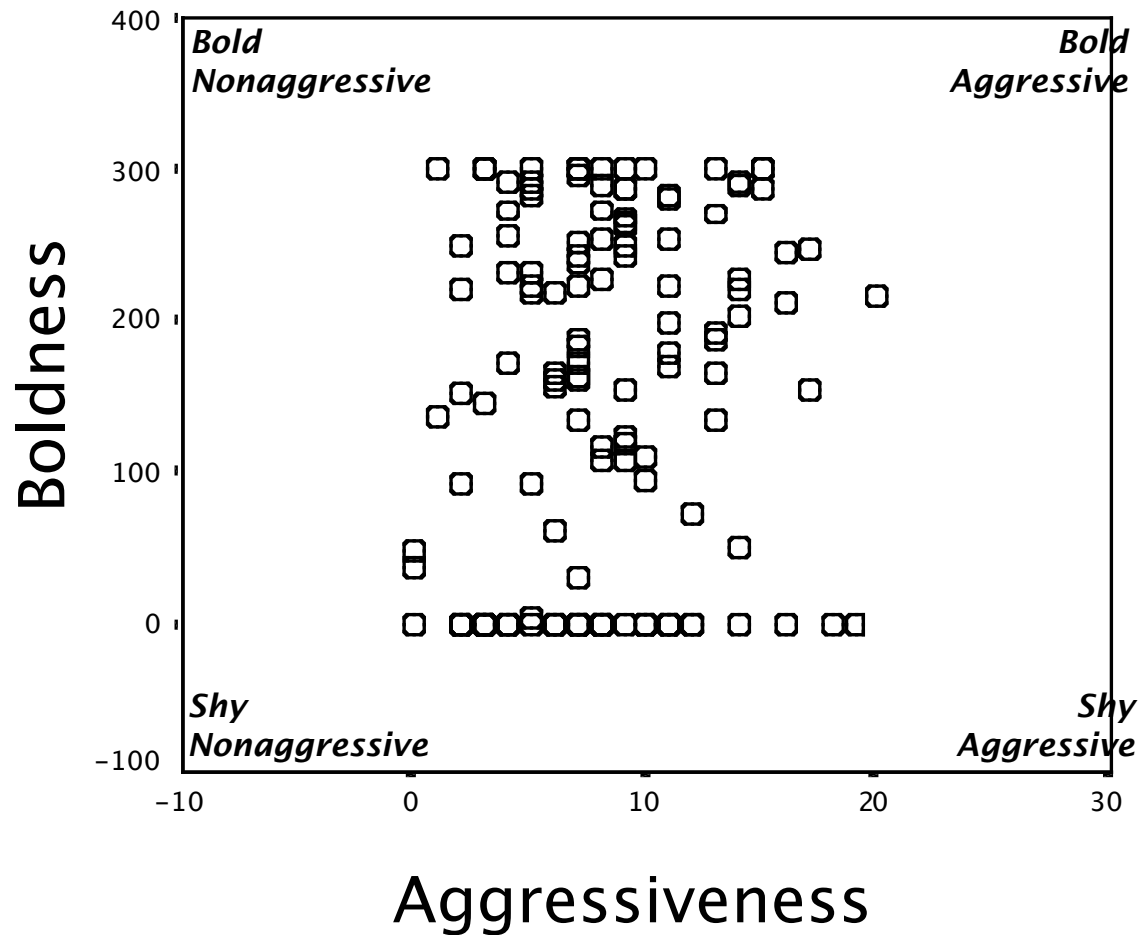
AFTER



Aggressiveness

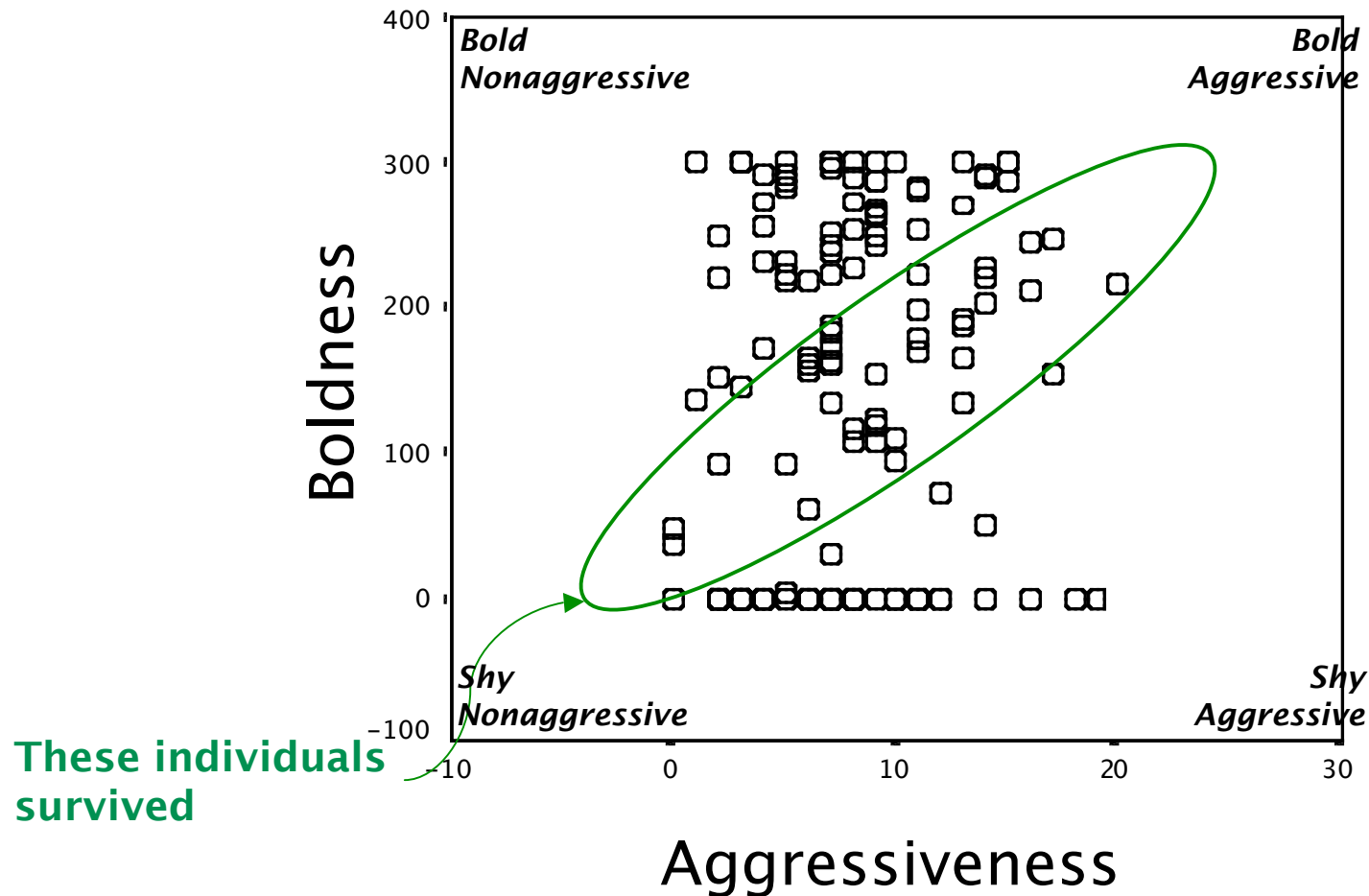
Control: Before NS, After NS
Tank: NS

Two ways this could have happened



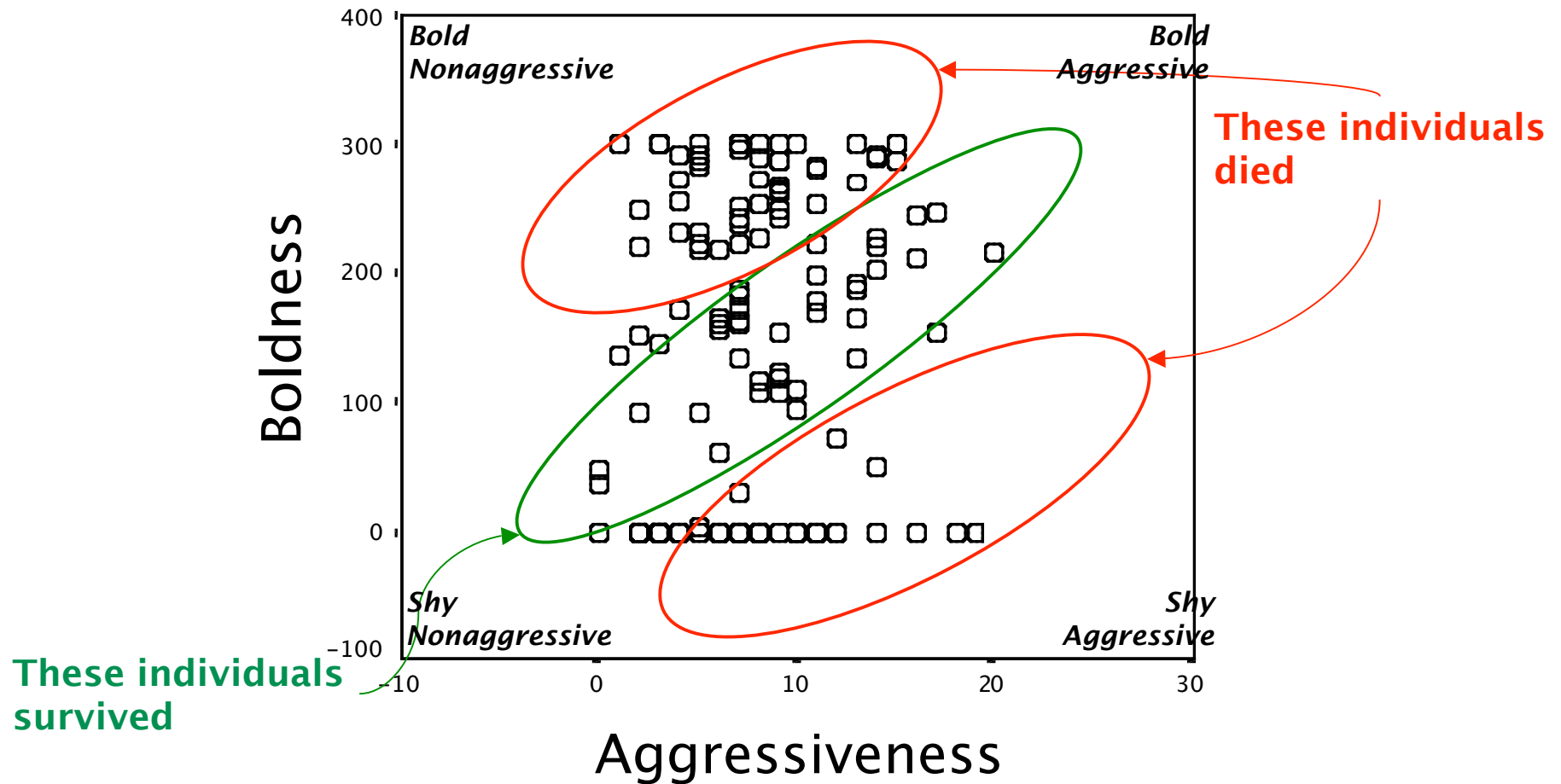
Two ways this could have happened

1. Selection via disproportionate survivorship



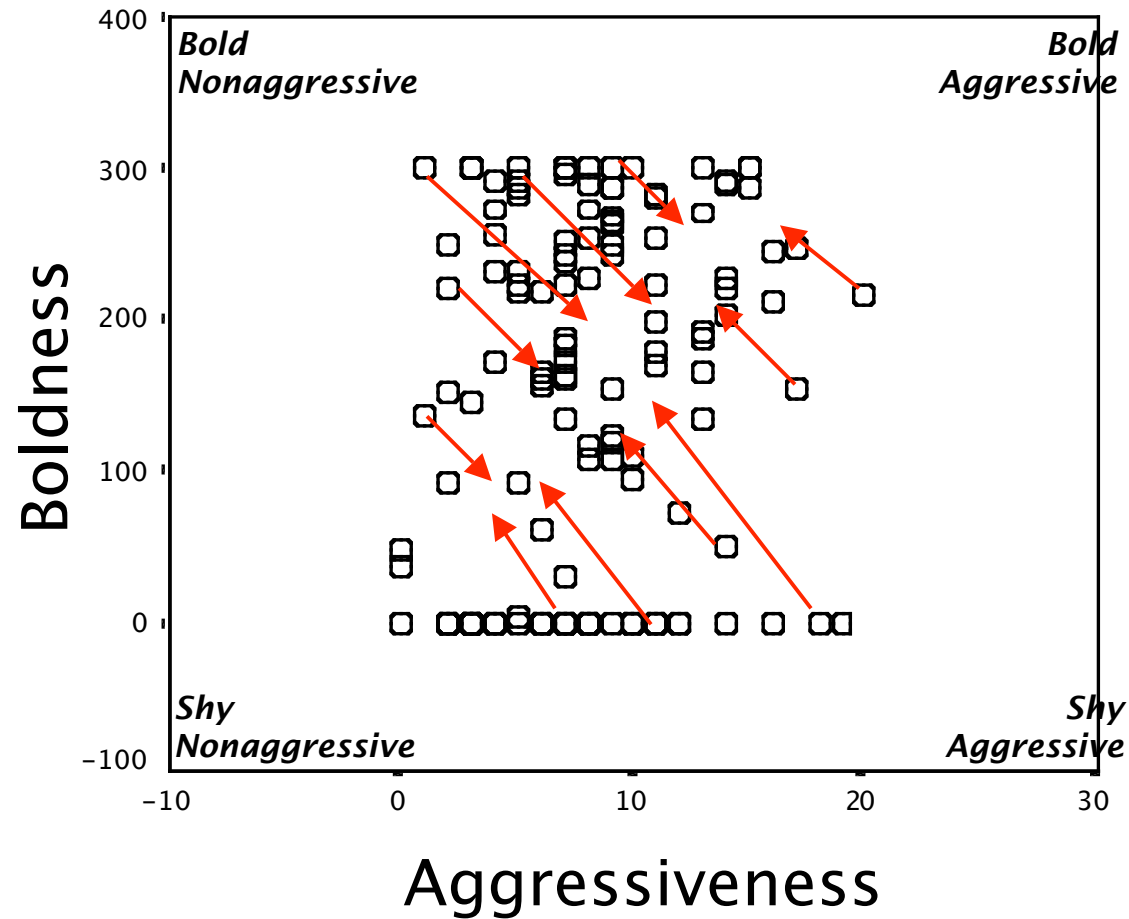
Two ways this could have happened

1. Selection via disproportionate survivorship



Two ways this could have happened

2. Behavioural plasticity

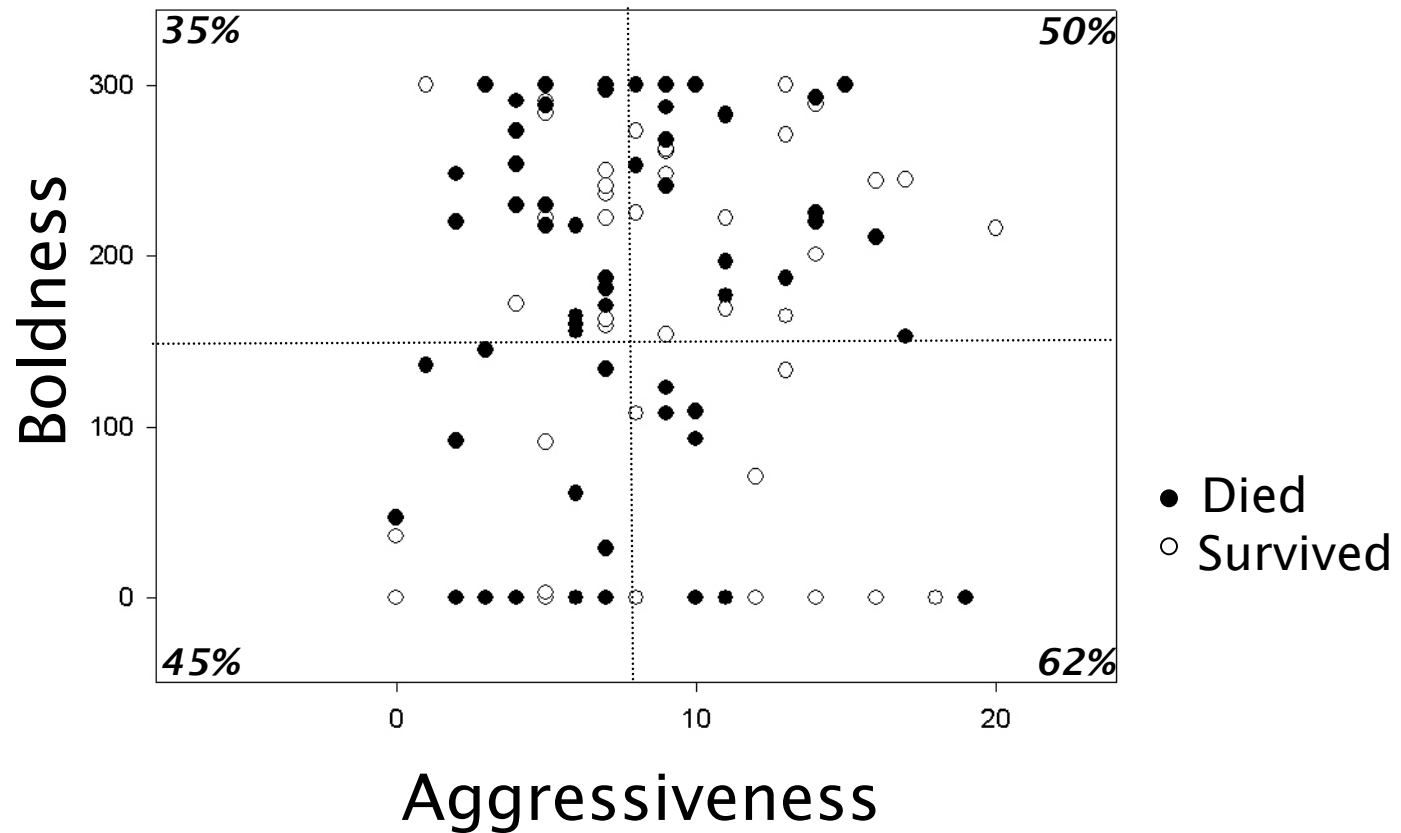


Directional selection favored increased aggressiveness, decreased boldness

Directional, stabilizing and correlational selection gradients ($\beta \pm$ standard error) from multiple logistic regression. Model $\chi^2 = 7.477$, 5 *df*, $n=122$, $p=0.188$.

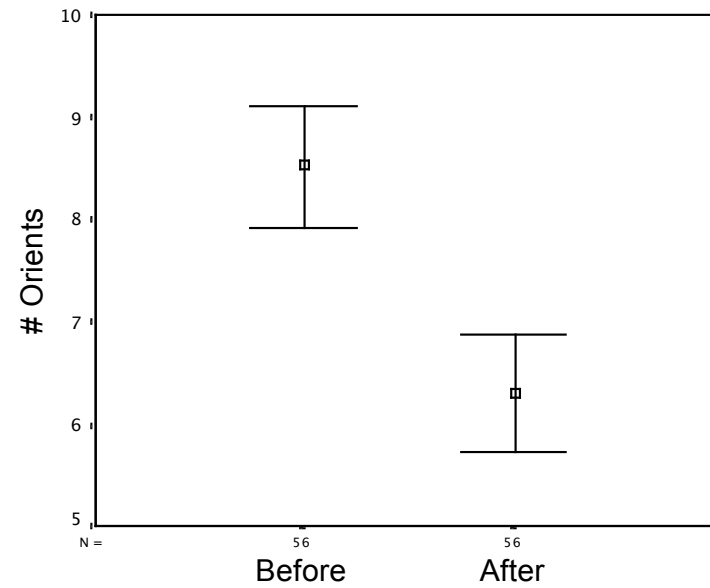
	$\beta \pm$ SE	<i>p</i>
Aggressiveness → # Orients	0.432 ± 0.212	0.04
Time eating	-0.330 ± 0.202	0.05
# Orients * Time eating	0.095 ± 0.193	0.62
Boldness → # Orients ²	-0.111 ± 0.156	0.47
Time eating ²	0.258 ± 0.294	0.38
Constant	-0.285 ± 0.368	0.43

Selection or plasticity?

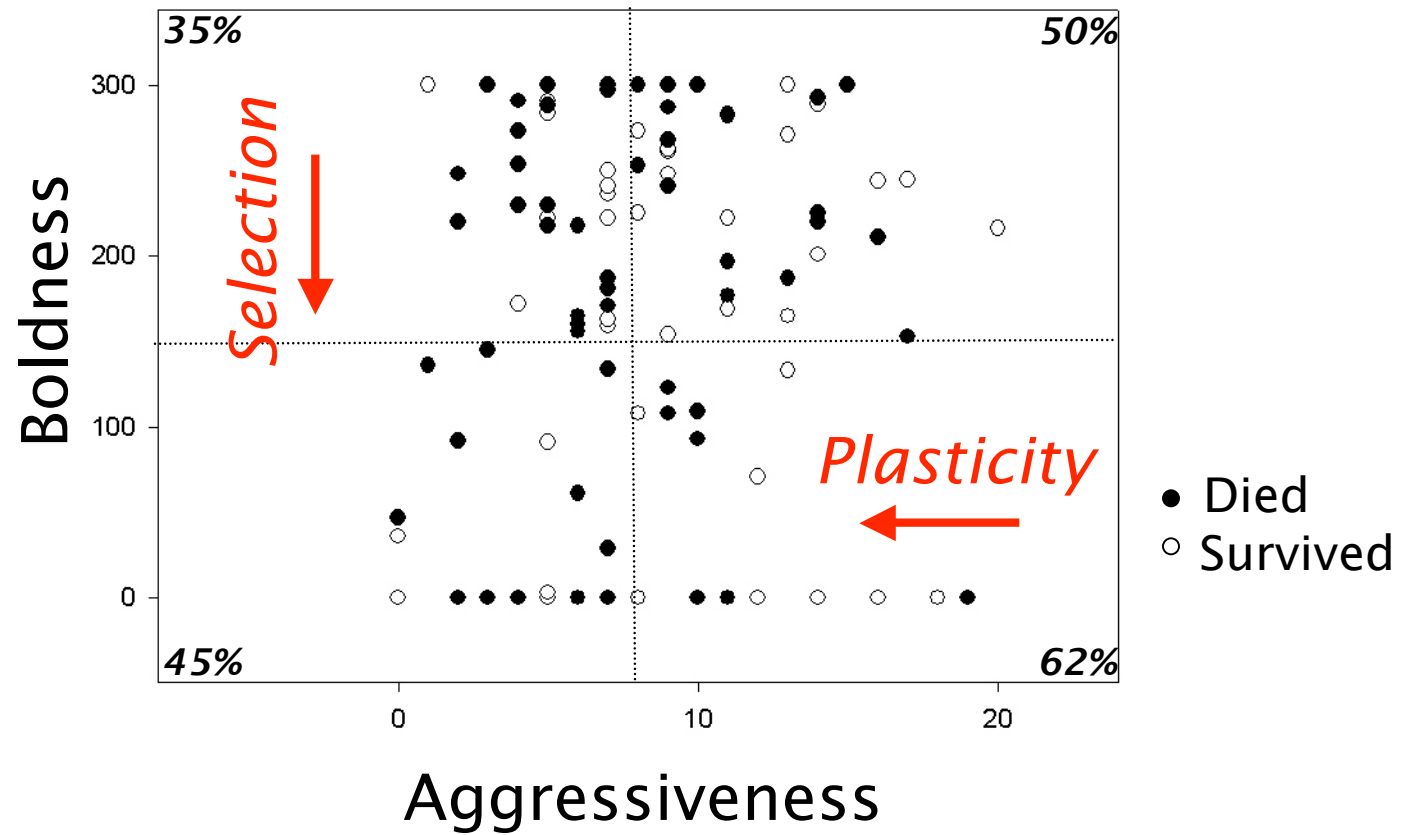


THERE WAS SELECTION, BUT IT WASN'T CORRELATIONAL

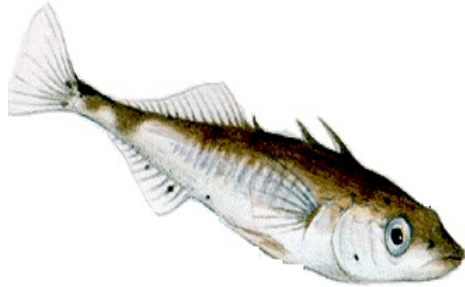
Sticklebacks became less aggressive



Selection and plasticity



Can experimental exposure to real predation risk generate the syndrome?



- Boldness and aggressiveness were not packaged together as a syndrome 'before' exposure to real predation risk.
- The boldness-aggressiveness behavioural syndrome appeared among the survivors.
- Both selection and plasticity generated the syndrome.

Can we make generalizations about the magnitude of consistent individual differences in behavior in non-human animals?

A repeatability estimate (the intraclass correlation coefficient) compares the extent of within versus between-individual variation.

We analyzed 759 repeatability estimates from 114 studies on 98 species.

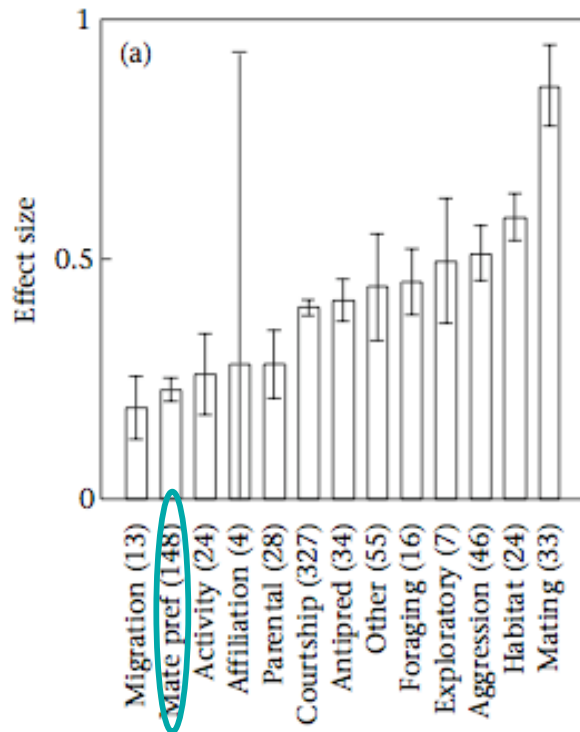
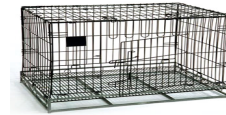
Factors: behaviour, taxa, age, number of measurements, field or lab study, sex



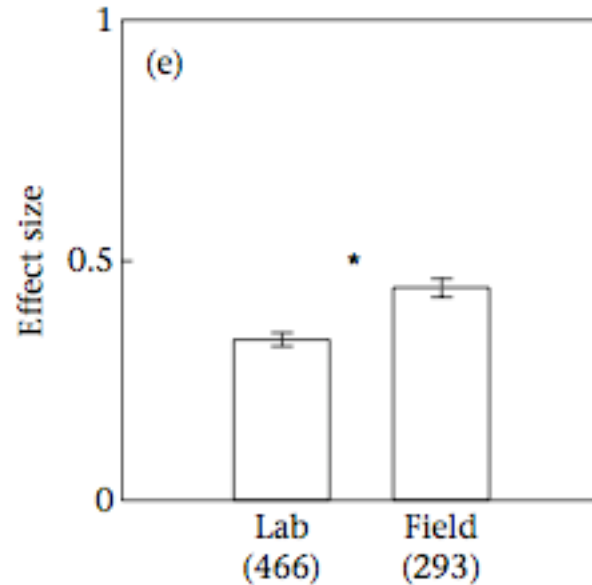
Bell, Hankison & Laskowski
2009 *Animal Behaviour*



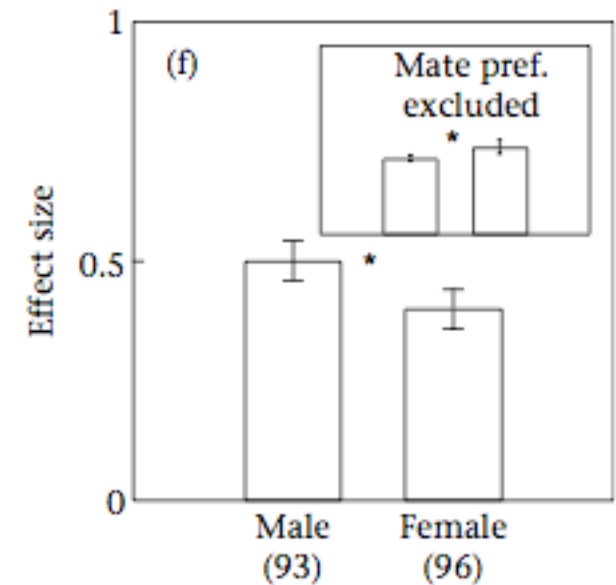
We can make some generalizations from the data



Some behaviours were more repeatable than others



Behaviours measured in the field were more repeatable than behaviours measured in the lab



Sex differences?

Some current projects in the lab....



Individual differences in response to a changing food resource (Kate Laskowski)

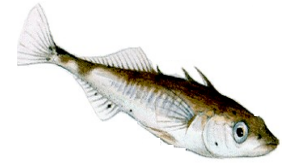
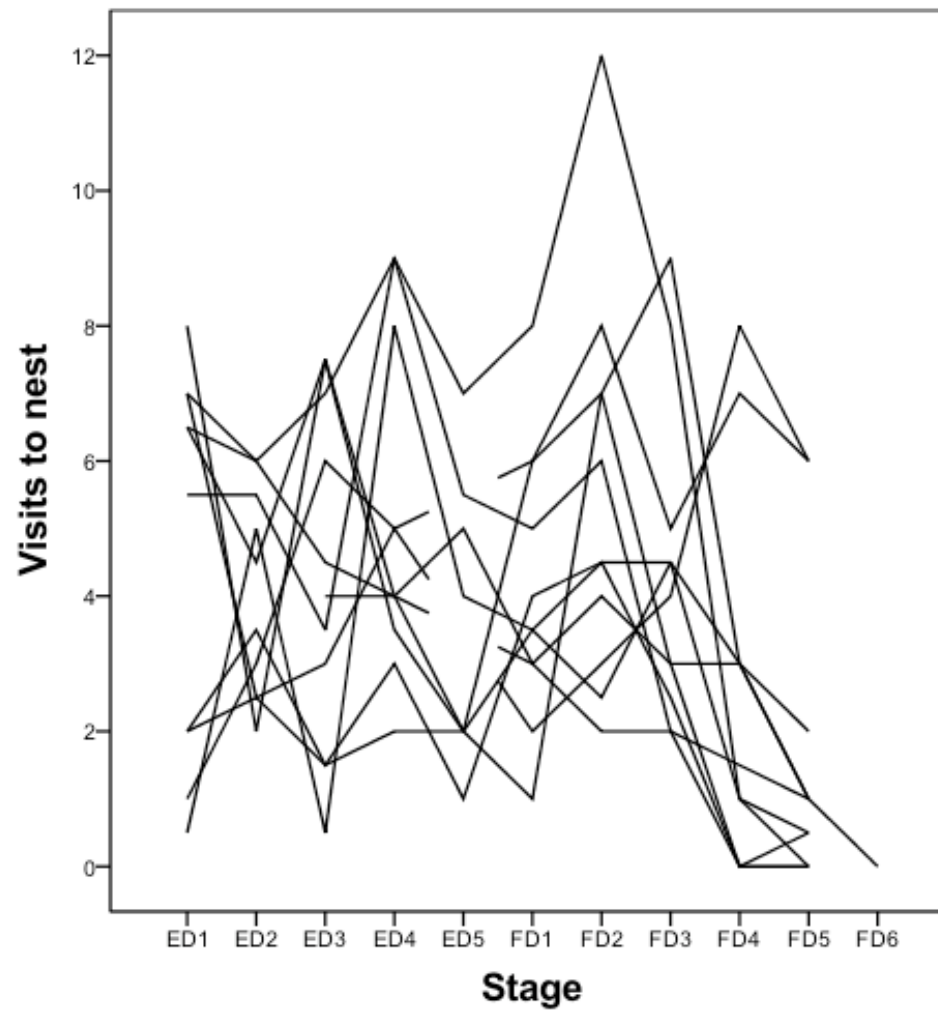


Individual differences in response to a changing food resource (Kate Laskowski)

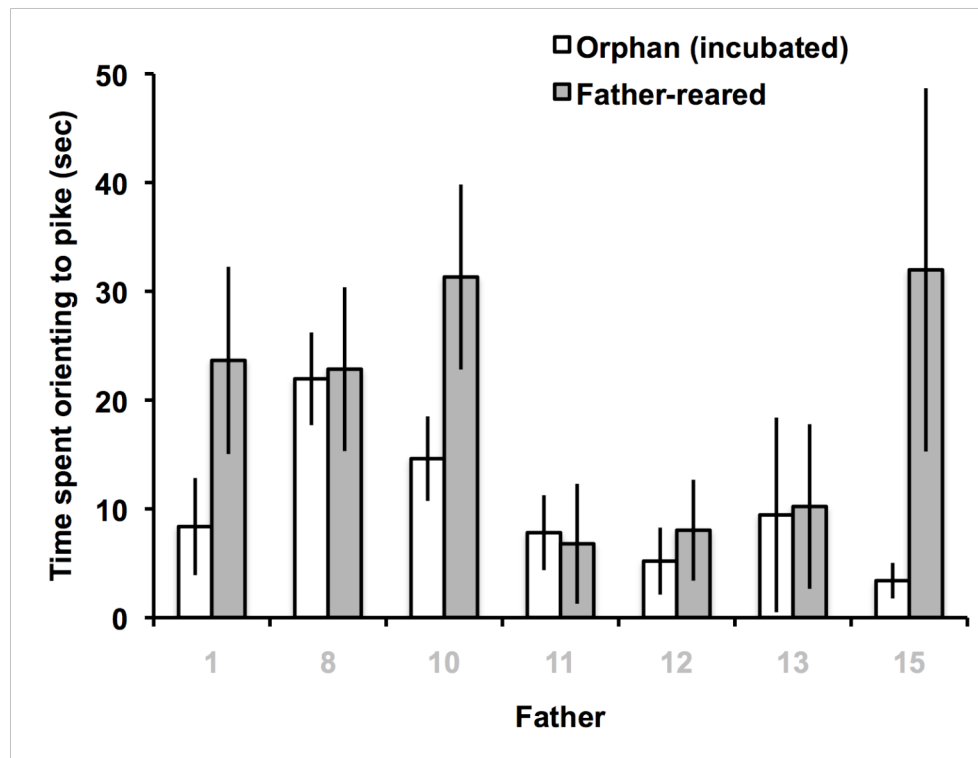
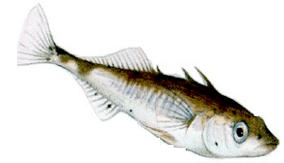


Individual differences in parenting behaviors (Laura Stein)

Individual differences in parenting behaviors (Laura Stein)



... and their consequences for offspring (Katie McGhee)

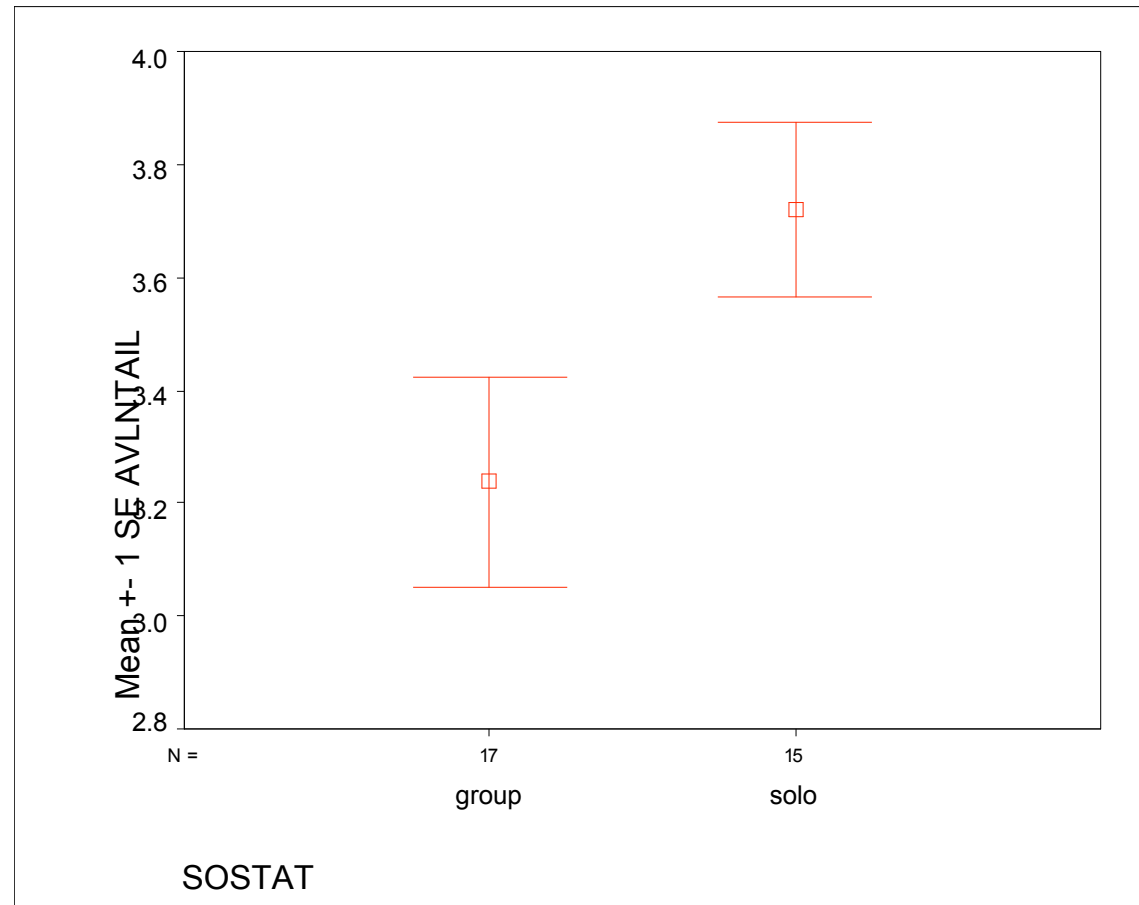


Individual differences in the latency to emerge and explore a novel environment *in the field* (Simon Pearish)

Individual differences in the latency to emerge and explore a novel environment *in the field* (Simon Pearish)

$r=0.67$, $n=32$
individuals
measured 3x

Individuals collected in groups emerged faster and explored more



Next: Diet?
Individual
specialization?

Searching the genome for the source of a behavioral syndrome in threespined sticklebacks

Alison M. Bell, Y. Osee Sanogo

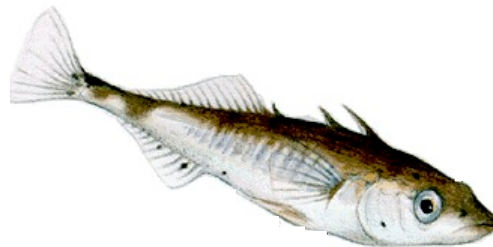
Integrative Biology, University of Illinois

Pengfei Yu, Dan Xie, Sheng Zhong

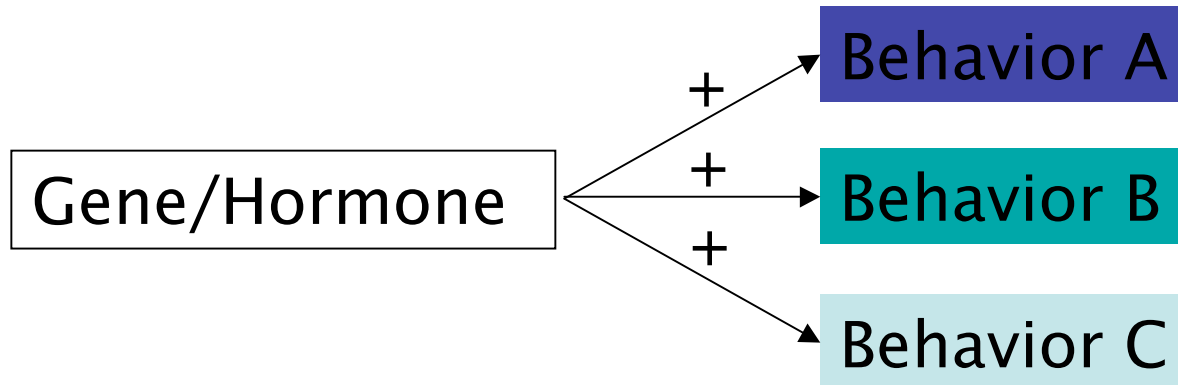
Bioengineering, University of Illinois

Mark Band

Keck Center, University of Illinois



Pleiotropy is the textbook cause of a behavioral syndrome



Aggressive behavior



Parental behavior

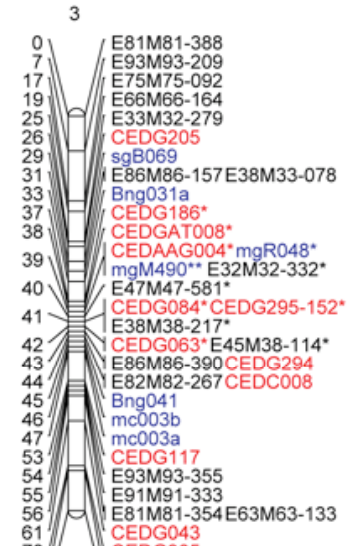
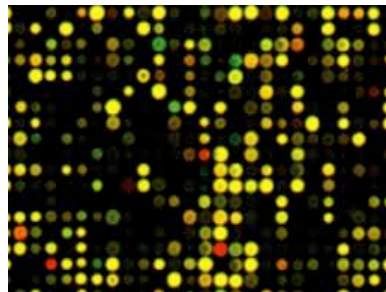
How to study pleiotropy?

- Quantitative genetics, genetic correlations

$$r_{Ax,y} = \frac{b_{xy}}{\sqrt{b_x b_y}}$$

- QTL mapping, gene association

- Whole genome expression



Hypothesis: Boldness toward predators and aggressiveness toward conspecifics are positively correlated because the same molecular mechanisms are involved in behavioral reactions to both predators and conspecifics.



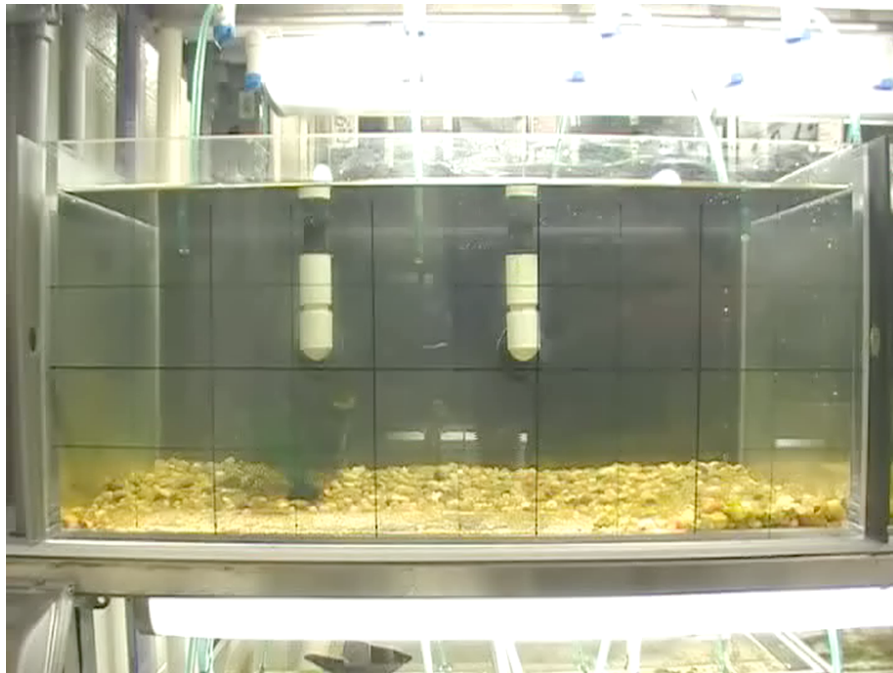
General strategy

- Use whole genome expression data to identify differentially expressed genes in response to **predators** and **conspecifics**.
- Look for genes that are shared.
- Search the genome for common regulatory motifs.



Methods: **Response to predators**

- Exposed sticklebacks to predator cues for six days
 - Control: no cues of predators
- mRNA from whole brain on day 7
- N=5 control, 5 experimental, 10 arrays
- 385K probe high density NimbleGen single-color microarray



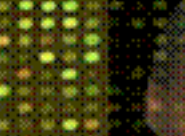
Methods: Response to a conspecific

- Exposed nesting male sticklebacks to a territorial intruder for 15 min
 - Control: empty flask
- Extracted mRNA from 4 brain regions 30 minutes after intruder removed
- N=5 control, n=4 experimental individuals, 40 arrays in loop design
- 44K probe, two-color Agilent microarrays



Are there any genes expressed in response to *predators* that also have a differential expression response to a *herbivore*?

AveExpr	t	P.Value	Name	logFC	AveExpr
6.80965806	7.71100353	2.00E-06	ENSGACT000C	1.53699956	8.55979
8.93053751	-6.26911607	1.99E-05	ENSGACT000C	1.46255835	8.45454
5.87794484	6.22591536	2.14E-05	ENSGACT000C	1.38815535	8.30059
5.65367305	6.16392829	2.88E-05	ENSGACT000C	1.73538125	8.44857
7.5873738	5.99099515	3.21E-05	ENSGACT000C	0.52885135	6.98584
6.7582394	5.85897362	4.03E-05	ENSGACT000C	0.62399932	7.01599
7.90325068	5.74846965	4.90E-05	ENSGACT000C	-0.71415381	11.2909
10.9963008	-5.71616589	5.18E-05	ENSGACT000C	2.03656931	7.94279
7.046863	-5.48148213	7.88E-05	ENSGACT000C	1.51117429	7.73648
6.60877028	5.44587923	8.40E-05	ENSGACT000C	0.68461468	8.77229



Osee Sanogo

expressed genes in response to predators

AveExpr	t	P.Value
6.80965806	7.71100353	2.00E-06
8.93053751	-6.26911607	1.99E-05
5.87794484	6.22591536	2.14E-05
5.65367305	6.16392829	2.88E-05
7.5873738	5.99099515	3.21E-05
6.7582394	5.85897362	4.03E-05
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10.9963008	-5.71616589	5.18E-05
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Differentially expressed genes in response to a predator

Name	logFC	AveExpr
ENSGACT000C	1.53699956	8.55979
ENSGACT000C	1.46255835	8.45454
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ENSGACT000C	-0.71415381	11.2909
ENSGACT000C	2.03656931	7.94279
ENSGACT000C	1.51117429	7.73648
ENSGACT000C	0.68461468	8.77229

Are there any genes on both lists?

Name	logFC	AveExpr	t	P.Value
ENSGACT000C	1.53699956	8.55975345	4.62764701	5.26E-05
ENSGACT000C	1.46255835	8.45454696	4.72708923	3.92E-05
ENSGACT000C	1.38815535	8.30059369	4.68097654	4.49E-05
ENSGACT000C	1.17353825	8.44857306	5.66347968	8.60E-06
ENSGACT000C	0.52885135	6.98584997	4.89515758	2.38E-05
ENSGACT000C	0.62399932	7.01595991	4.52349969	7.14E-05
ENSGACT000C	-0.71415381	11.2909507	-4.63982281	5.07E-05
ENSGACT000C	2.03655931	7.94275716	5.80702091	1.57E-06
ENSGACT000C	1.51147429	7.73648503	6.74560726	9.73E-08
ENSGACT000C	-0.68484168	8.77225459	-4.84728232	2.75E-05
ENSGACT000C	0.28205416	6.94793683	5.03528721	4.08E-05
ENSGACT000C	-0.75393092	7.70287065	-4.35904739	0.00011563
ENSGACT000C	-1.07371357	7.73734707	-5.35410858	1.84E-05
ENSGACT000C	-0.77885333	6.9476131	-5.91263028	4.68E-06
ENSGACT000C	-0.7428225	7.33317776	-4.71229381	9.17E-05
ENSGACT000C	-0.63735243	7.80541663	-4.29425443	0.00013965
ENSGACT000C	-1.72459378	10.4241775	-4.93919214	2.09E-05
ENSGACT000C	-1.51201562	10.7788351	-4.56648403	6.30E-05
ENSGACT000C	-0.75914885	6.92851184	-5.78316616	1.68E-06
ENSGACT000C	-0.67543919	6.92804603	-4.18144716	0.00019371
ENSGACT000C	1.66679674	9.37491742	5.14375697	3.11E-05
ENSGACT000C	0.83530847	13.4748094	4.72761741	3.91E-05
ENSGACT000C	1.00349112	13.2062186	4.25138774	0.00015818
ENSGACT000C	-1.34685001	7.86247332	-5.28038768	2.22E-05
ENSGACT000C	-1.0983276	11.8291236	-4.54210242	6.76E-05
ENSGACT000C	-0.88736034	11.4406665	-4.31096401	0.00013302
ENSGACT000C	0.6656206	8.29392013	4.17018873	0.00020012
ENSGACT000C	0.50211368	8.66918733	4.42506912	9.53E-05
ENSGACT000C	0.38353182	9.78453306	4.22244662	0.00017203
ENSGACT000C	-1.17449125	8.72349206	-4.52948619	0.0001451
ENSGACT000C	-0.44789614	8.34775216	-4.99854484	4.47E-05
ENSGACT000C	0.59757929	9.84414604	5.00200698	4.43E-05
ENSGACT000C	0.58569949	8.13500543	4.17236049	0.00019887
ENSGACT000C	0.63086789	9.27170965	4.2674784	0.00015095
ENSGACT000C	0.58722091	8.01995173	4.17459693	0.00019758
ENSGACT000C	-1.5469682	7.37196514	-5.38540332	1.71E-05
ENSGACT000C	-1.22689152	6.90958149	-5.73872075	7.15E-06
ENSGACT000C	-0.68166903	6.94858525	-5.63630045	2.61E-06
ENSGACT000C	-0.46580229	7.03483345	-4.31263304	0.00013238
ENSGACT000C	-0.76386438	7.19029991	-5.94174115	1.05E-06
ENSGACT000C	-0.92557084	7.43980589	-6.00562874	8.68E-07
ENSGACT000C	-0.83628694	7.72481566	-5.16231387	1.08E-05
ENSGACT000C	-0.99624592	7.63446231	-5.28766443	7.40E-06
ENSGACT000C	0.70138835	8.63219863	4.72706112	8.83E-05
ENSGACT000C	-0.56300685	7.67575255	-5.44748485	4.59E-06

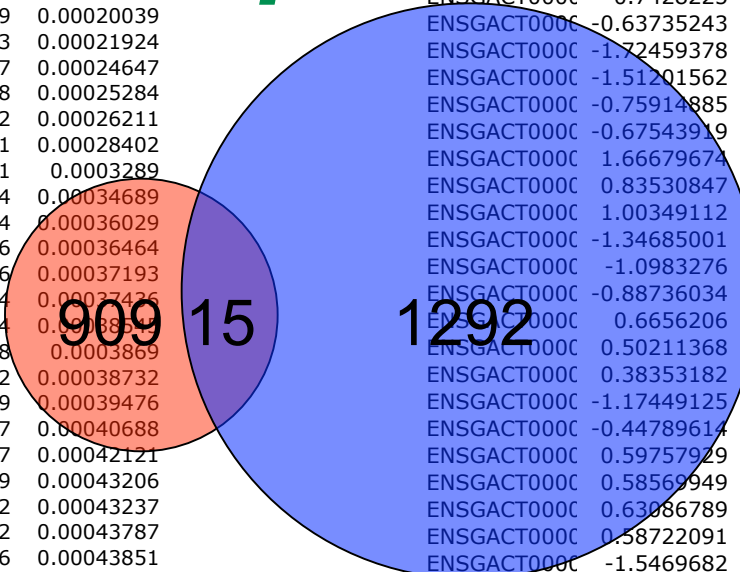
Differentially expressed genes in
response to a *predator*

Differentially expressed genes in
response to a *conspecific*

ID	logFC	AveExpr	t	P.Value
ENSGACT000C	3.96400072	6.80965806	7.71100353	2.00E-06
ENSGACT000C	-1.71111924	8.97053751	-6.26911607	1.99E-05
ENSGACT000C	1.27767463	5.87794474	6.25915366	2.1E-05
ENSGACT000C	1.47331778	5.55373355	5.16392829	2.3E-05
ENSGACT000C	-1.82268859	7.5873738	-5.99099515	3.21E-05
ENSGACT000C	2.61609287	6.7582394	5.85897362	4.03E-05
ENSGACT000C	2.9095063	7.90325068	5.74846965	4.90E-05
ENSGACT000C	-1.28103675	10.9963008	-5.71619381	5.18E-05
ENSGACT000C	-1.45118577	7.046863	-5.4814121	5.88E-05
ENSGACT000C	1.06389949	6.60877028	5.44587923	8.40E-05
ENSGACT000C	-1.04482182	7.83176392	-5.40878232	8.98E-05
ENSGACT000C	-1.15796511	5.21660596	-5.40700538	9.01E-05
ENSGACT000C	-1.25706822	7.90217103	-5.36477497	9.73E-05
ENSGACT000C	-1.43729735	10.2231468	-5.29313488	0.00011086
ENSGACT000C	1.17374329	9.60978592	5.1922135	0.00013333
ENSGACT000C	1.05453454	8.14411738	5.16103391	0.00014124
ENSGACT000C	1.05604658	10.4382316	4.9724209	0.00020039
ENSGACT000C	-1.07846171	7.59630601	-4.92435043	0.00021924
ENSGACT000C	0.96311602	11.4608616	4.86194777	0.00024647
ENSGACT000C	-1.77662012	6.99994263	-4.84837968	0.00025284
ENSGACT000C	-1.22630355	6.95061188	-4.82925672	0.00026211
ENSGACT000C	1.18014254	9.16346671	4.78670981	0.00028402
ENSGACT000C	-1.79854069	10.5330242	-4.7092631	0.0003289
ENSGACT000C	-1.47308447	7.25243051	-4.68123254	0.00034689
ENSGACT000C	1.38980207	9.08624235	4.66131544	0.00036029
ENSGACT000C	0.96704301	8.56739755	4.65501676	0.00036464
ENSGACT000C	-1.28621653	5.42774887	-4.64463586	0.00037193
ENSGACT000C	1.21975293	7.22679384	4.64122004	0.00037426
ENSGACT000C	1.7233546	11.7544607	4.62591384	0.00038547
ENSGACT000C	-0.96649046	5.55239457	-4.62394378	0.0003869
ENSGACT000C	1.18801288	5.35343288	4.62336862	0.00038732
ENSGACT000C	-1.14871759	5.45177113	-4.61340469	0.00039476
ENSGACT000C	1.42495573	8.22709944	4.59758087	0.00040688
ENSGACT000C	1.04471947	5.7608261	4.57947767	0.00042121
ENSGACT000C	1.06904589	6.47656381	4.56619109	0.00043206
ENSGACT000C	1.30521539	6.94758214	4.56581402	0.00043237
ENSGACT000C	0.93537996	6.55639315	4.55922372	0.00043787
ENSGACT000C	0.91413447	6.47630261	4.55846026	0.00043851
ENSGACT000C	-1.12913415	6.68136437	-4.54401583	0.00045082
ENSGACT000C	1.38258806	9.20343883	4.53824474	0.00045583
ENSGACT000C	1.21428229	6.36987912	4.53395674	0.0004596
ENSGACT000C	0.91870473	6.73292996	4.51312734	0.00047835
ENSGACT000C	1.14304683	7.7299686	4.5116584	0.0004797
ENSGACT000C	1.13953425	5.14171384	4.50236383	0.00048834
ENSGACT000C	1.33234962	7.66072363	4.49208818	0.00049809
ENSGACT000C	1.12315127	5.57353927	4.48110326	0.00050872

Name	logFC	AveExpr	t	P.Value
ENSGACT000C	1.53699956	8.55975345	4.62764701	5.26E-05
ENSGACT000C	1.46255935	8.45454696	4.72708923	3.92E-05
ENSGACT000C	1.33855955	8.3659369	4.68097654	4.49E-05
ENSGACT000C	1.77353825	8.44897306	5.66347968	8.60E-06
ENSGACT000C	0.52885135	6.98584997	4.89515758	2.38E-05
ENSGACT000C	0.62200932	7.01595991	4.52349969	7.14E-05
ENSGACT000C	1.71419381	11.2909507	-4.63982281	5.07E-05
ENSGACT000C	2.3656931	7.94275716	5.80702091	1.57E-06
ENSGACT000C	1.51117429	7.73648503	6.74560726	9.73E-08
ENSGACT000C	-0.68461468	8.77225459	-4.84728232	2.75E-05
ENSGACT000C	0.28205416	6.94793683	5.03528721	4.08E-05
ENSGACT000C	-0.75393092	7.70287065	-4.35904739	0.00011563
ENSGACT000C	-1.07371357	7.73734707	-5.35410858	1.84E-05
ENSGACT000C	-0.77885333	6.9476131	-5.91263028	4.68E-06
ENSGACT000C	-0.7428225	7.33317776	-4.71229381	9.17E-05
ENSGACT000C	-0.63735243	7.80541663	-4.29425443	0.00013965
ENSGACT000C	-1.72459378	10.4241775	-4.93919214	2.09E-05
ENSGACT000C	-1.51201562	10.7788351	-4.56648403	6.30E-05
ENSGACT000C	-0.75914885	6.92851184	-5.78316616	1.68E-06
ENSGACT000C	-0.67543919	6.92804603	-4.18144716	0.00019371
ENSGACT000C	1.66679674	9.37491742	5.14375697	3.11E-05
ENSGACT000C	0.83530847	13.4748094	4.72761741	3.91E-05
ENSGACT000C	1.00349112	13.2062186	4.25138774	0.00015818
ENSGACT000C	-1.34685001	7.86247332	-5.28038768	2.22E-05
ENSGACT000C	-1.0983276	11.8291236	-4.54210242	6.76E-05
ENSGACT000C	-0.88736034	11.4406665	-4.31096401	0.00013302
ENSGACT000C	0.6656206	8.29392013	4.17018873	0.00020012
ENSGACT000C	0.50211368	8.66918733	4.42506912	9.53E-05
ENSGACT000C	0.38353182	9.78453306	4.22244662	0.00017203
ENSGACT000C	-1.17449125	8.72349206	-4.52948619	0.0001451
ENSGACT000C	-0.44789614	8.34775216	-4.99854484	4.47E-05
ENSGACT000C	0.59757929	9.84414604	5.00200698	4.43E-05
ENSGACT000C	0.58569949	8.13500543	4.17236049	0.00019887
ENSGACT000C	0.63086789	9.27170965	4.2674784	0.00015095
ENSGACT000C	0.58722091	8.01995173	4.17459693	0.00019758
ENSGACT000C	-1.5469682	7.37196514	-5.38540332	1.71E-05
ENSGACT000C	-1.22689152	6.90958149	-5.73872075	7.15E-06
ENSGACT000C	-0.68166903	6.94858525	-5.63630045	2.61E-06
ENSGACT000C	-0.46580229	7.03483345	-4.31263304	0.00013238
ENSGACT000C	-0.76386438	7.19029991	-5.94174115	1.05E-06
ENSGACT000C	-0.92557084	7.43980589	-6.00562874	8.68E-07
ENSGACT000C	-0.83628694	7.72481566	-5.16231387	1.08E-05
ENSGACT000C	-0.99624592	7.63446231	-5.28766443	7.40E-06
ENSGACT000C	0.70138835	8.63219863	4.72706112	8.83E-05
ENSGACT000C	-0.56300685	7.67575255	-5.44748485	4.59E-06

Are there any genes
on both lists?
A few



Transcription factors (TFs)

- TFs regulate the expression of other genes by binding to the promoter region of a gene.
- The TF binding site sequence is conserved.
- We searched the genome upstream (1500bp) of our *candidate genes* to look for binding sites of 570 TFs, and compared them to the promoter regions of *other genes*.



Dan Xie, Bioengineering



Transfac, Prima V1.0
UCSC Genome Browser
(Stickleback, gasAcu1)

We identified TF motifs for both up- and down-regulated genes

Predator	Conspecific		
Up (24)	Up (52)		
Down (26)	Down (22)		

Some motifs were common to responses to both predators and a conspecific

Predator	Conspecific	Overlapping motifs	Transcription factors
Down (26)	Up (52)	2	ETS, OCT-1
Up (24)	Up (52)	4	AP-4, MCM1, Pax-6, OCT-1
Down (26)	Down (22)	2	HSF, OCT-1
Up (24)	Down (22)	2	Skn-1, OCT-1

What's next?

Compare 'baseline' gene expression between different behavioral types of individuals or populations.

Hypothesis: The same set of genes discriminates between shy vs bold individuals/populations, and nonaggressive vs aggressive individuals/populations

