The yellow gene regulates behavioural plasticity by repressing male courtship in *Bicyclus anynana* butterflies

Heidi Connahs1,†, Eunice Jingmei Tan1,2,†, Yi Ting Ter1,†, Emilie Dion1, Yuji Matsuoka1, Ashley Bear3 and Antónia Monteiro1,2

1Department of Biological Sciences, National University of Singapore, 14 Science Drive 4, Singapore 117543
2Yale-NUS College, 16 College Avenue West, Singapore 138527
3Department of Ecology and Evolutionary Biology, Yale University, CT 06511, USA

†Indicate equal contributions.

Seasonal plasticity in male courtship in *Bicyclus anynana* butterflies is due to variation in levels of the steroid hormone 20E (20-hydroxyecdysone) during pupation. Wet season (WS) males have high levels of 20E and become active courtiers. Dry season (DS) males have lower levels of 20E and reduced courtship rates. However, WS courtship rates can be achieved if DS male pupae are injected with 20E at 30% of pupation. Here, we investigated the genes involved in male courtship plasticity and examined whether 20E plays an organizational role in the pupal brain that later influences the sexual behaviour of adults. We show that DS pupal brains have a sevenfold upregulation of the yellow gene relative to the WS brains, and that knocking out yellow leads to increased male courtship. We find that injecting 20E into DS pupa reduced yellow expression although not significantly. Our results show that yellow is a repressor of the neural circuity for male courtship behaviour in *B. anynana*. 20E levels experienced during pupation could play an organizational role during pupal brain development by regulating yellow expression, however, other factors might also be involved. Our findings are in striking contrast to *Drosophila* where yellow is required for male courtship.

1. Introduction

Behavioural plasticity is essential for animals to adapt to environmental variation and it is often triggered by hormonal changes that organize or activate neural circuits in the brain [1–3]. Seasonal changes in temperature and photoperiod can serve as important cues that alter hormone levels and sexual behaviour in a wide range of vertebrate and invertebrate taxa [4]. Precisely how hormone signalling influences sexual behaviour and the identity of the downstream genes in most animals however, is not well known [5]. In vertebrates, hormones are considered to play both a brain organizational role during development, as well as a behavioural activational role in adults, compared to just an adult activational role in insects [6,7]. Whether hormones also display a similar latent role in insects remains unclear [7].

Most research on the genetics of courtship behaviour is based on studies in *Drosophila* [8,9]. Among the most well-studied genes are the transcription factors *fruitless* (*fru*) and *doublesex* (*dsx*), which undergo alternative splicing to produce male-specific proteins [10]. One of the genes proposed to be downstream of *fru* is *yellow*, a gene that is unique to insects and appears to have numerous biological functions, including melanin pigmentation, cuticle formation, butterfly scale colour and morphology and also male sexual behaviour [9,11–15]. Exactly how yellow expression influences male courtship behaviour in *Drosophila* has been an important topic of investigation that has yielded conflicting results.
Early studies suggested that yellow male mutants were less successful during courtship due to neural circuitry regulation of courtship song [8,9,16]. More recent work demonstrated that yellow does not influence male courtship through any neural circuitry but rather by influencing the mechanical structure of the male sex combs [15]. These findings, therefore, contradict the long-held assumption that yellow is part of the neural circuitry governed by

$fri^{M}$ that regulates male courtship behaviour.

Sexual behaviour in insects has traditionally been viewed as a consequence of cell-autonomous processes taking place during brain development, and involving sex determination genes [6] such as fruitless and doublesex [17]. The role of insect hormones is typically described as playing an activating role, allowing rapid and reversible behavioural changes, such as activating neural circuits that regulate pheromone communication or sexual receptivity [18–21]. However, hormones have also been proposed to play an organizational role in insects, for instance in the regulation of behavioural polyphenisms in honeybees and locusts [7] and sexual maturity in Drosophila [22]. Yet, no evidence is available for the organizational role of steroid hormones in driving sexual behaviours in adult insects, similar to the latent role of steroid hormones in vertebrate sexual differentiation, where exposure to different hormone levels during ontogeny leads to discrete, fixed differences in neural development and sexual behaviours [7].

For insects living in seasonal environments, hormones could play an organizational role earlier in development to ensure that sexual behaviour is optimized for particular environmental conditions that will be prevalent upon adult emergence [23]. An example of a species where such a brain organizational role may be happening is the African seasonal polyphenic butterfly, Bicyclus anynana. This species shows an interesting sex-role reversal between seasonal forms that develop at different temperatures, and where temperature cues in the arrival of different seasons. In particular, wet season (WS) males, reared at high temperatures, play the active courting role, while dry season (DS) males, reared at low temperatures court less and become the choosy sex [24]. The adaptive reason driving courtship plasticity in males is associated with increased reproductive costs for DS males, which provide females with beneficial spermatophores [24]. Provision of this spermatophore ultimately shortens DS male lifespan, but lengthens DS female lifespan and helps them survive through the more stressful and resource-limited DS [23,24]. While the behavioural ecology of these butterflies may explain seasonal variation in male courtship rates, the neural re-wiring that switches the male behaviour is completely unknown.

In B. anynana, signalling of the hormone 20-hydroxyecdysone (20E) during early pupal development has been shown to regulate male courtship [25]. Throughout pupal development, there are significantly lower levels of 20E titres circulating in the haemolymph of DS than of WS males [25]. The reduced courtship of DS males, however, can be switched to the WS active courting form by rearing pupae at higher temperatures during the first 50% of pupal development [23] or, alternatively, by keeping the pupae at low temperatures but injecting them with 20E at 30% of pupal development [25]. These experiments suggest that this pupal stage is a critical period that determines male sexual behaviour and that 20E may play an organizing role in the developing male brain of B. anynana. However, we currently have no direct evidence that 20E acts specifically on the brain to alter expression of genes that regulate male courtship behaviour in adults. High levels of 20E could upregulate genes required for active courtship such as those described for Drosophila including fruitless, doublesex and yellow [9,15,17]. Alternatively, 20E signalling may not lead to appreciable differences in the neural circuitry of dry and WS male brains, but may instead influence other phenotypic traits that are also important in courtship behaviour such as pheromone production [26] or the UV brightness of eyespot centres [24].

We investigated whether genes are differentially expressed in DS and WS male brains at 30% of pupal development, and which genes, if any, are responding to 20E injections at that same time period. We conducted a transcriptome analysis on dissected brains of vehicle-injected DS and WS forms, and brains of DS forms injected with 20E, all at 2 h after the injection. Our transcriptome analysis revealed that the yellow gene is differentially expressed between seasonal forms. We then pursued additional qPCR experiments and functional studies by knocking out yellow using CRISPR-Cas9 and observing the courtship behaviour of mutant and wild-type males.

2. Material and methods

(a) Transcriptome assembly and analyses

To mimic DS and WS conditions, caterpillars of B. anynana (established in New Haven, CT, from an original Malawi stock collected in 1988) were reared under WS and DS temperatures in climate-controlled rooms at 27°C and 17°C, respectively, at 80% humidity, and a 12 : 12 h light : dark photoperiod. Caterpillars were fed corn plants ad libitum until pupation. Pupae were staged, such that the per cent of pupal development was known for all individuals. At 30% of pupal development (day 2 in the WS butterflies and day 6 in the DS butterflies) DS pupae were injected with either 3 µl of 2000 pg µl$^{-1}$ (6000 pg total) (10% 20E in EtOH + 90% saline) of 20E (Sigma-Aldrich) or with 3 µl of vehicle (10% EtOH and 90% saline) and WS pupae were injected with 3 µl of vehicle in the lateral posterior region of the fifth abdominal segment. The injections were done at 12.00 h and the brains were dissected 2 h later, at 14.00 h. We injected the pupae 2 h before brain collection because previous studies have demonstrated that genes, which respond early to 20E signalling, are expressed about 2 h after exposure to 20E [27,28].

Each sample consisted of three biological replicates of DS pupal brains following treatment with vehicle only (WSV); four biological replicates of DS pupal brains treated with vehicle only (DSV); and five biological replicates of DS pupal brains treated with 20E (DS20E). Each biological replicate contained the brains of five male pupae. We assembled the transcriptome using 12 RNA-Seq libraries. Details of the sample preparation and transcriptome assembly can be found in the electronic supplementary material, Methods section.

The differentially expressed genes (DEGs) were annotated with Blast2GO v. 5.2.5. We used the public NCBI Blast service (QBlast) to blast our sequences against the non-redundant protein database using the blastx-fast program. Matched transcripts were filtered using a cut-off E-value of $1 \times 10^{-3}$; otherwise the default settings for Blast2GO were used at each step. To annotate the remaining transcriptome, we performed a local blastx of the assembled contigs against the B. anynana v. 1.2 draft genome [29].

(b) qPCR sample collection and experiments

Sample collection was similar to that described above for the RNA-Seq experiment (see electronic supplementary material, Methods
section and table S1). At 30% development, we injected pupae with 20E or vehicle solutions using the same protocol as described above. Pupae were kept in their respective development temperatures and brains were dissected 2 h, 4 h and 24 h after injections in ice cold 1× PBS, placed immediately into RNALater (Qiagen, GmbH, Hilden, Germany) and stored at −20°C until RNA extraction. We used five biological replicates per treatment, each made of five pooled brains (for the 2 h dissections) or two pooled brains (for the 4- and 24-hr dissections).

(c) Behavioural assays
We conducted behavioural assays using a Yellow mutant line to investigate whether loss of yellow function affects butterfly courtship behaviour. To establish the Yellow mutant line, we inserted an attP sequence into exon 4 of the yellow gene to disrupt its overall sequence (electronic supplementary material, figure S1). We used a knock-in method through homology-directed repair (HDR) using a single-stranded DNA (ssDNA) as a template. The ssDNA construct was made following methods described in [30]. Further details are in the electronic supplementary material, Methods section.

Behavioural assays were conducted in cylindrical hanging cages (30 × 40 cm) under one full-spectrum light tube (Plantmax) and one UV light bulb (Arcadia Marine Blue), at 23°C, from 17.00 to 18.00. This specific time of observation was chosen because B. anynana exhibits crepuscular courtship [25]. Visual barriers were placed between cages to prevent mate-copying [31]. Within each sex, butterflies used for each assay were of the same age. All butterflies used in the assays ranged from 4 to 8 days old. Two experiments were performed, one with live females and the other with decapitated live females. The treatments were (i) two Wt males × two Wt females and (ii) two Yellow mutant males × two Wt females (electronic supplementary material, figure S2). One of the two males/females in an assay was dotted with a black marker at both of its ventral hindwings to allow for sex-specific scoring of behaviour. The multiple elements of courtship, as documented in Nieberding et al. [32], were scored in the assays: (1) localization (flying to other butterfly), (2) rapid flickering of wings, (3) thrusting (touching female’s wings with head) and (4) attempting (curling of the abdomen). Orientation (orienting body to female’s posterior) was not recorded since it was difficult to score or interpret their intent (courtship or coincidence) with that behaviour. Latency to mate (time taken from the start of the assay to the first mating) and mating duration were recorded as well. Behavioural assays lasted 1 h, and quantification of an individual male’s behaviour stopped once it mated but the other male’s behaviour was still quantified until its own mating, or 1 h, had elapsed.

For decapitation assays, only females were decapitated to characterize male sexual behaviour in the absence of female response [33]. Decapitated females were first anaesthetized in a −20°C freezer for 20 min and their heads were removed. Females were pinned through their thorax into opposite sides of the cage as illustrated in electronic supplementary material, figure S2c,d. The same behavioural scoring as described above was used for the decapitation assays as well.

(d) Statistical analysis
The data were evaluated for equality of variances and normality using the Levene’s test and Shapiro–Wilk test, respectively. Total duration and frequency of courtship were calculated by adding up the duration/ frequency of all the courtship elements displayed (localizing + flicker + thrust + attempt) during the observation period. A generalized linear model (GLM) with a Tweedie distribution (gamma family; tweedie package [34]) was done to test the impact of the treatment type (Yellow/Wt) and season (WS/DS) on the total duration of courtship. A Tweedie distribution (gamma) was used due to the high number of zeroes and the skewness of data. The impact of the treatment type (Yellow/Wt) and season (WS/DS) on the frequency of each courtship element was compared using a zero-inflated Poisson model with a negative binomial distribution to account for overdispersion (pscl package) [35]. Both mating latency and mating duration of the mated pairs were compared using independent t-tests. χ²-tests were carried out to identify any associations between treatment type and mating success. Statistical tests and figures were done with IBM SPSS Statistics 25 and R-v. 4.0.2 [36]. The spectral data of eyespots were visualized using the pavo package (electronic supplementary material, figure S3 [37]). Spectral analysis was done through calculating area under curve (AUC) for each eyespot replicate and the AUC analysed using an ANOVA with post hoc Tukey test in R.

3. Results
The critical window for determining male courtship behaviour occurs at 30% of pupation, as both temperature shift experiments or injections of 20E into DS pupa at this specific stage, rescues WS courtship rates [25]. Therefore, we reasoned that temperature may be affecting 20E levels, and 20E may be affecting gene expression in the pupal brain causing a switch in adult behaviour from a choosy male to an active courter. To test this hypothesis, we compared the transcriptome of DS male brains injected with 20E at 30% pupal development with DS and WS male brains injected with a vehicle solution at the same developmental stage (DSV and WSV, respectively) to identify genes that may be regulating courtship behaviour (figure 1).

We identified 399 DEGs between DSV and WSV pupal brains, 302 were upregulated and 97 were downregulated in DSV. Comparing DS20E pupal brains to WSV, we identified 399 DEGs, 291 were upregulated and 108 were downregulated in DS20E. Comparing DS20E pupal brains to DSV we identified 151 DEGs, 79 were upregulated and 72 were downregulated in DS20E. Overall, the smaller number of DEGs observed between DSV and DS20E (151) compared to DSV and WSV (399) suggest that the DS20E brain transcriptome profile was more similar to DSV than WSV at 2 h post-injection (electronic supplementary material, figure S4). Full summary statistics are provided in electronic supplementary material, table S2 and figures S4–S5. The list of DEGs can be found in electronic supplementary material, tables S3–S4.

(a) The yellow gene which drives male courtship in
_Drosophila_ was significantly upregulated in DSV and DS20E pupal brains
The melanin pathway gene _yellow_ was significantly upregulated in both DSV (sevenfold increase) and DS20E (eightfold increase) when compared to WSV. Other genes of interest pertaining to courtship behaviour included genes involved in the Juvenile hormone signalling pathway, such as _Juvenile hormone esterase_ (JHE), which was downregulated in DSV compared to WSV, and _juvenile hormone epoxide hydrolase-like_ (JHEH, which hydrolyses JH) which was upregulated in DSV and DS20E when compared to WSV. We also identified a gene involved in dopamine metabolism, _dopamine N-acetyltransferase_ (AANAT1/ DAT1) which was upregulated in DS20E (but not in DSV) when compared to WSV. Genes involved in neural development included _neuropeptide CCHamide2_, _Neural Wiskott–Aldrich_
Importance for male courtship in Drosophila melanogaster is an interesting candidate to explore further due to its effect in the dry season (DS) versus the wet season (WS) courtship rates. Yellow protein is downregulated in D20E compared to WSV. (a) Genes were not differentially expressed at 30% of pupal development. (b) Overview of the experimental set-up for the pupal injections, brain dissections, and RNA-seq for the three different treatment groups. (Online version in colour.)

(b) 20E downregulates the expression of yellow 4 h after injections

Yellow is an interesting candidate to explore further due to its importance for male courtship in Drosophila [9,15]. As the transcriptome showed that it was significantly upregulated in both DS treatments compared to WS, we considered whether the 2 h time point for brain dissections was too early to detect any effects of 20E on yellow expression. We hypothesized that injection of 20E into DS male pupae would reduce yellow expression down to WS levels at a later time point post-injection. Similar to the transcriptome experiment, we injected 20E in DS male pupae at 30% development [25] and a vehicle solution in both DS and WS pupae at the same stage, and used qPCR to...
measure expression of yellow in dissected pupal brains at 2, 4 and 24 h post injection.

Two hours post-injection, yellow expression was 2.5 times higher in pupal brains of both DSV and DS20E compared to the expression in WSV pupal brains, (mirroring our RNA-seq results), although the expression levels were not significantly different (electronic supplementary material, figure S6, ANOVA: F = 0.50, p = 0.62). At 4 h post-injection, expression of yellow increased significantly by eightfold in DSV compared to WSV pupal brains (figure 3, ANOVA: F = 5.43, p = 0.023; post hoc analysis WSV-DSV: adj. p = 0.027), while the level of yellow expression in DS20E remained low, similar to those of WSV brains (ANOVA post hoc analysis WSV-DS20E: adj. p = 0.79). At 24 h post-injection, expression levels of yellow in the pupal brains were 2.8 (DSV) and 3.7 (DS20E) times higher than in WSV (electronic supplementary material, figure S6). Relative levels of yellow expression were significantly higher in DS20E than in WSV pupal brains (ANOVA: F = 4.12, p = 0.046; post hoc analysis WSV versus DS20E: adj. p = 0.046). These results demonstrate that a single injection of 20E into DS pupae at 30% of development was sufficient to decrease yellow expression levels in DS20E to WSV levels at 4 h post-injection. This single injection did not impact yellow levels at the earlier 2 h time period, nor kept yellow levels low at 24 h post injection, suggesting that a short interval of time around 30% pupal development encompasses a hormone-sensitive window in which yellow is downregulated by 20E to WS levels.

(c) Yellow males courted more frequently and for a longer duration than Wt males

Based on the transcriptome and qPCR results we hypothesized that yellow is a repressor of male courtship as DS males exhibit lower courtship than WS males and have significantly higher expression of yellow during pupal brain development. Using a Yellow mutant line (hereafter Yellow males), we compared the duration and frequency of the Yellow males and the wild-type (Wt) male courtship sequence, including copulation, in both seasonal forms. These behavioural assays were initially performed with live females. Yellow males courted for a longer duration (WS: t = 2.181, p = 0.0323; DS: t = 2.083, p = 0.0416; figure 4a) than Wt males regardless of seasonality. Yellow males also courted more frequently (WS: z = 2.465, p = 0.0137; figure 4b) than Wt males, but this was only observed in the WS form. In addition, Yellow males remained in copulation longer with live females (t = 2.174, p = 0.039; figure 4c) than Wt males, but this was only observed in the DS form.

We found that both Yellow and Wt males had similar ultraviolet (UV) reflectivity in their dorsal eyespots— a known sexual ornament (electronic supplementary material, figure S3) [39,40]. Thus, females would not be choosing males based on their dorsal UV reflective eyespots. Yellow males displayed an overall lighter pigmentation compared to Wt males, which might influence female’s receptivity to mating with a novel phenotype. Hence, we repeated these male courtship observations using freshly decapitated females. Decapitation prevents important visual cues detected by a female from impacting a male’s behaviour, such as more intense courtship provoked by a female’s increased rejection behaviour [33]. Yellow males still courted for a longer duration (WS: t = 2.269, p = 0.0266; DS: t = 3.21,

(d) Yellow males copulated longer in their DS form than WS form

To test whether differences in yellow expression levels were sufficient to explain courtship differences between the seasonal forms, we compared the duration and frequency of courtship between DS and WS Yellow males. If Yellow, alone, was responsible for courtship differences between the forms, then DS and WS Yellow males should display similar levels of courtship. There were no differences in the duration or frequency of courtship between DS and WS Yellow males (figures 4a and b). However, there was a significant difference in mating duration between the Yellow seasonal forms. With live females, DS Yellow males copulated longer (t = −2.119, p = 0.0425; figure 4c) than WS Yellow males. Similar behaviour was observed in males courting decapitated females, with DS Yellow males copulating longer (t = −2.34, p = 0.0318; figure 4c) than their WS counterparts.

4. Discussion

In insects, hormones are typically assumed to regulate sexual behaviour by activating existing neural circuits that control processes such as sexual maturation, memory formation and pheromone communication [19,41–43]. It remains unclear, however, whether insect hormones also exhibit a latency effect whereby exposure earlier in development can organize neural circuits that affect sexual behaviour in adults. The identity of these hormone-sensitive genes involved in neural circuits regulating male courtship also remains unknown. Although, we did not find conclusive evidence that 20E plays
an organizational role, we show that yellow is involved in regulating seasonal courtship plasticity by repressing male courtship behaviour.

(a) Yellow functions as a repressor of male courtship in B. anynana

We show, using RNA-seq, that yellow is significantly upregulated by sevenfold in the pupal brains of DS male butterflies which court less than WS males. This increase in yellow expression could be in response to seasonal fluctuations in 20E, as injection of this hormone into DS males at 30% of pupal development rescues WS courtship levels [25]. Our qPCR results for 4 h post-injection of 20E into DS pupa, showed that while levels of yellow expression declined to similar levels observed in WS males, the difference was not significant from DS levels. High levels of yellow in DS males suggested that yellow was a repressor of courtship. This was confirmed by knocking out yellow in B. anynana and observing Yellow males exhibiting increased courtship frequency and

Figure 4. Courtship behaviour of Wt and Yellow males, for wet (WS) and dry (DS) seasonal forms, in both live and decapitation assays. Yellow males courted at a higher duration and frequency than Wt males, for both WS and DS forms. DS Yellow males remained in copula for a longer period of time than WS Yellow males. (a) Courtship duration, (b) courtship frequency and (c) mating duration were quantified. Mating duration was quantified among mated males only. Vertical bars represent mean ± SEM. Open circles are data points. Asterisks indicate significant differences: *p ≤ 0.05, **p ≤ 0.01, ***p ≤ 0.001. Outliers are removed from the figures (Q1 – 3*IQR or Q3 + 3*IQR). n(WS-Live-Wt) and n(WS-Live-Yellow) = 38, n(DS-Live-Wt) and n(DS-Live-Yellow) = 34, n(DS-Decap-Wt) and n(DS-Decap-Yellow) = 30. (Online version in colour.)
duration compared to Wt males of both seasonal forms. Given that Yellow WS males displayed more active courtship than Wt WS males, this suggests that low levels of yellow expression are still required in Wt WS males to reduce courtship and optimize energy expenditure, as increased wing fluttering observed in Yellow males did not translate to increased mating success (electronic supplementary material, figure S7).

Our findings are in striking contrast to those observed in *Drosophila* where yellow is required for normal male courtship behaviour and male mating success. However, in *B. anynana* males, yellow functions as a repressor of male courtship and does not disrupt mating success. Early studies in *Drosophila* suggested that yellow mutant males were less active during courtship and that mutations in yellow disrupted wing extension during the courtship ritual, preventing males from performing a courtship song which is required for male mating success [9,10]. However, recent work by Massey et al. argued that a lack of melanization in the sex combs of yellow mutants, rather than any impairment in neural circuitry affecting courtship song, was the trait that prevented males from successfully grasping females [16]. All research to date on yellow mutants in *Drosophila*, however, clearly demonstrates that yellow is absolutely required for successful male courtship and mating success.

In *B. anynana*, it is possible that changes in yellow expression in tissues other than the brain might have influenced courtship behaviour of Yellow males. As butterflies do not have sex combs we can rule out this possibility. In the Yellow males, the white centre of the eyespot, which is crucial for courtship behaviour shows no difference in brightness between Yellow and Wt males (electronic supplementary material, figure S3) [39,40]. Thus, we conclude that our findings point toward yellow influencing neural circuits regulating behaviour.

In our experimental set-up, using two males and two females, we found no difference in courtship rates between DS and WS wild-type butterflies. This result conflicts with previous work showing that WS males court more than DS males in larger social groups [24,25]. This difference may be explained by the reduction of sexual competitors in our assay, previously shown to increase the likelihood of courting and copulating by DS males in this species [44].

(b) Yellow may influence courtship behaviour in *B. anynana* via the dopaminergic signalling pathway

Our findings raise questions about how yellow, a gene involved in melanization, might regulate courtship behaviour. Melanin synthesis enzymes are expressed in the *Drosophila* brain and may be involved in the production of neuromelanin in dopaminergic neurons [45]. Yellow is thought to function as a dopachrome conversion enzyme (DCE) in the melanin pathway converting L-Dopa to Dopamine [11,12]. L-Dopa is also used as a substrate for dopamine which is involved in both cuticle pigmentation and neurotransmission [46]. Dopaminergic signalling has been shown to regulate mating drive and persistence and duration of mating in male *Drosophila* [46,47]. Thus, variation in yellow expression could alter the availability of L-Dopa for dopamine synthesis, with higher expression of yellow in DS brains leading to reduced L-Dopa. Alternatively, Yellow may physically bind to dopamine, as demonstrated in a study of salivary proteins in sandflies [48]. Thus, increased expression of yellow in DS brains could lead to reduced dopamine availability, which may inhibit courtship behaviour. In electronic supplementary material, figure S8, we suggest a possible mechanism of Yellow involvement in the pathway converting tyrosine to L-Dopa in dopaminergic neurons.

Currently, we have no direct evidence that dopamine levels differ between DS and WS *Bicyclus* brains. However, a few genes from our transcriptome analyses may provide some indirect evidence. We found that dopamine N-acetyltransferase, AANAT/DAT1 was upregulated in pupal brains of DS20E (but not in DSV) as compared to WSV brains. This may indicate a transient response to the 20E injection. The function of AANAT is to metabolize and inactivate secreted dopamine in the synapse shortly after release [45,49]. In young female *Drosophila virilis*, higher titres of 20E lead to an increase in dopamine, although this appears to be associated with reduced activity of AANAT [50]. However, in retinal cells of fish, AANAT activity is positively correlated with dopamine levels [51]. An increase in dopamine induced by 20E would provide a mechanistic explanation as to why DS male pupa injected with 20E display active WS courtship behaviour.

We also observed changes in Juvenile hormone (JH) signalling, which is known to interact with dopamine to affect sexual maturity and courtship behaviour in *Drosophila*, likely through changes in neural development [22,52]. In DS pupal brains, JHE was downregulated and Juvenile hormone epoxide hydrolase (JHEH), which degrades JH [53], was upregulated. These findings could indicate that dopamine levels are low in DS male pupal brains as dopamine increases JH titres in young female *D. virilis* by inhibiting its degradation [54]. JH is also associated with increased dopamine levels in male honeybees [55]. Although we also see an upregulation of JHEH but no longer a downregulation of JHE in DS20E, this may reflect a response to changing levels of dopamine. Interactions between 20E, JH, dopamine and AANAT in *Drosophila* represent a complex pathway as depicted in [50,56] thus, we must interpret our findings with caution. However, given that this is an important pathway for regulating courtship behaviour in *Drosophila*, differential expression of these genes in our transcriptome analyses suggest their possible involvement in regulating courtship behaviour in *B. anynana*. Surprisingly, genes important for male courtship in *Drosophila* (*fruitless, dsx crol, lola, cadN and chimno*) [38], all showed very low levels of expression and were not differentially expressed in *B. anynana* male brains (figure 2).

5. Conclusion

Here, we provide the first evidence for a gene regulating butterfly courtship behaviour. In *B. anynana*, yellow functions as a repressor of male courtship thus higher expression of this gene in DS pupal brains may explain why these butterflies court less than WS males. It remains unclear, however, why injection of 20E into DS pupae at 30% development converts their behaviour to the WS form. Further work is required to confirm whether 20E plays a role in organizing neural circuits (potentially involving yellow) during critical windows of brain development. Our results also suggest possible interactions of 20E on JH and dopamine signalling, a circuit well described in *Drosophila*. Future studies examining dopamine levels between the seasonal forms and the individual role of 20E and JH on dopaminergic signalling would help clarify
mechanistically why the yellow gene functions as a repressor of male courtship in these butterflies. For animals living in seasonal environments, selection may favour adaptations that use external cues to optimize behaviour, such as employing environmentally induced hormones, even in insects.

Ethics. The research reported in this paper, which involved insects only, was exempt from ethics approval procedures by the National University of Singapore.

Data accessibility. Raw reads of the RNA-seq libraries were uploaded to the SRA database with the SRA accession no. PRJNA544388. This Transcriptome Shotgun Assembly project has been deposited at DDBJ/EMBL/GenBank under the accession no. GHRJ0000000. The annotated transcriptome is available from the Dryad Digital Repository: https://doi.org/10.5061/dryad.9cnp9hqg4 [57].

The data are provided in electronic supplementary material [58].

Authors’ contributions. H.C.: conceptualization, formal analysis, investigation, visualization, writing—original draft, writing—review and editing; E.J.T.: conceptualization, formal analysis, investigation, visualization, writing—original draft, writing—review and editing; Y.T.T.: conceptualization, formal analysis, investigation, visualization, writing—original draft, writing—review and editing; E.D.: conceptualization, formal analysis, investigation, visualization, writing—original draft, writing—review and editing; Y.M.: conceptualization, investigation, visualization, writing—original draft, writing—review and editing; A.B.: conceptualization, investigation, A.M.: conceptualization, funding acquisition, project administration, resources, writing—original draft, writing—review and editing.

All authors gave final approval for publication and agreed to be held accountable for the work performed therein.

Competing interests. The authors declare no competing interests.

Funding. E.J.T. was supported by a Postdoctoral Fellowship from Yale-NUS College. We thank NSF award no. DDIG IOS-1110523 to A.M. and A.B., Ministry of Education Singapore, award no. MOE2018-T2-1-092 to A.M. and National Research Foundation, Singapore, Investigatorship award no. NRF-NRFIF05-2019-0006 to A.M.

Acknowledgements. We would like to acknowledge Dong Qiang Cheng, for helping E.J.T. with the hardware issues on the laboratory server; Jan Gruber, for advising E.J.T. regarding the Linux operating system and also Firefly Farms in Singapore for supplying corn for larval rearing.

References

1. Neal JK, Wade J. 2007 Courtship and copulation in the adult male green anole: effects of season, hormone and female contact on reproductive behavior and morphology. Behav. Brain Res. 177, 177–185. (doi:10.1016/j.bbr.2006.11.035)
2. Meitzenn J, Thompson CK. 2008 Seasonal-like growth and regression of the avian song control system: neural and behavioral plasticity in adult male Gambel’s white-crowned sparrows. Gen. Comp. Endocrinol. 157, 259–265. (doi:10.1016/j.ygcen.2008.03.014)
3. Forfano PM, Sinnero JA, Rohmann KN, Bass AH. 2015 Neuroendocrine control of seasonal plasticity in the auditory and vocal systems of fish. Front. Neuroendocrinol. 37, 129–145. (doi:10.1016/j.ynefn.2014.08.002)
4. Tramontin AD, Brenowitz EA. 2000 Seasonal plasticity in the adult brain. Trends Neurosci. 23, 251–258. (doi:10.1016/S0166-2236(00)15158-7)
5. Zhang SX, Glantz EH, Miner LE, Rogulja D, Crickmore AD. Temporal and spatial expression of the yellow gene in correlation with cuticle formation and DOPA decarboxylase activity in Drosophila development. Dev. Biol. 129, 1849–1858. (doi:10.1242/dev.129.8.1849)
6. Walter MF, Black BC, Afsar H, Kermabon AJ, Wright TRF, Biessmann H. 1991 Seasonal and spatial expression of the yellow gene in correlation with cuticle formation and DOPA decarboxylase activity in Drosophila development. Dev. Biol. 147, 32–45. (doi:10.1016/0022-1669(90)90004-3)
7. Matsuoka Y, Monteiro A. 2018 Melanin pathway genes regulate color and morphology of butterfly wing scales. Cell Rep. 24, 56–65. (doi:10.1016/j.celrep.2018.05.092)
8. Massey JH, Chung D, Siwanowicz I, Stern DL, Yamamoto D. 2008 Fruitless and doublesex genes regulate color and morphology of butterfly wing scales. Developmental plasticity in sexual roles of butterfly aglottis iphila. Dev. Biol. 317, 1849–1858. (doi:10.1242/dev.129.8.1849)
9. Walter MF, Black BC, Afsar H, Kermabon AJ, Wright TRF, Biessmann H. 1991 Seasonal and spatial expression of the yellow gene in correlation with cuticle formation and DOPA decarboxylase activity in Drosophila development. Dev. Biol. 147, 32–45. (doi:10.1016/0022-1669(90)90004-3)
10. Matsuoka Y, Monteiro A. 2018 Melanin pathway genes regulate color and morphology of butterfly wing scales. Cell Rep. 24, 56–65. (doi:10.1016/j.celrep.2018.05.092)
11. Massey JH, Chung D, Siwanowicz I, Stern DL, Yamamoto D. 2008 Fruitless and doublesex genes regulate color and morphology of butterfly wing scales. Cell Rep. 24, 56–65. (doi:10.1016/j.celrep.2018.05.092)
12. Bear A, Monteiro A. 2013 Male courtship rate plasticity in the butterfly Bicyclus anynana is controlled by temperature experienced during the pupal and adult stages. PLoS ONE 8, e64061. (doi:10.1371/journal.pone.0064061)
13. Prudic KL, Jean C, Cao H, Monteiro A. 2011 Developmental plasticity in sexual roles of butterfly species drives mutual sexual ornamentation. Science 331, 73–75. (doi:10.1126/science.1197114)
14. Bear A, Prudic KL, Monteiro A. 2017 Steroid hormone signaling during development has a latent effect on adult male sexual behavior in the butterfly Bicyclus anynana. PLoS ONE 12, e0174403. (doi:10.1371/journal.pone.0174403)
15. Dion E, Monteiro A, Yew JY. 2016 Phenotypic plasticity in sex pheromone production in Bicyclus anynana butterflies. Sci. Rep. 6, 39002. (doi:10.1038/srep39002)
16. Karim FD, Guild GM, Thummel CS. 1993 The Drosophila broad-complex plays a key role in controlling ecdysone-regulated gene expression at...
the onset of metamorphosis. Development 118, 977–988. (doi:10.1242/dev.118.3.977)

28. Gauhar Z, Sun W, Hua S, Mason CE, Fucho F, Li TR, Boutros M, White KP. 2009 Genomic mapping of binding regions for the Ecdysone receptor protein complex. Genome Res. 19, 1006–1013. (doi:10.1101/gr.18349.108)

29. Newell RW et al. 2017 A high-coverage draft genome of the mycaleus butterfly Bicyclus anynana. Gigascience 6, 1–7. (doi:10.1093/gigascience/gix035)

30. Miura H, Quadros RM, Gunemruth CB, Ohtsuka M. 2018 Easi-CRISPR for creating knock-in and conditional knockout mouse models using long ssDNA donors. Nat. Protoc. 13, 195–215. (doi:10.1038/nprot.2017.153)

31. Merry F, Varela SAM, Danchin E, Blanchet S, Panjoo D, Coolen I, Wagner RH. 2009 Public versus personal information for mate copying in an invertebrate. Curr. Biol. 19, 730–734. (doi:10.1016/j.cub.2009.02.064)

32. Nieberding CM et al. 2008 The male sex pheromone of the butterfly Bicyclus anynana: towards an evolutionary analysis. PLoS ONE 3, e2751. (doi:10.1371/journal.pone.0002751)

33. Speth HF. 1966 Drosophilid mating behaviour: the behaviour of decapitated females. Anim. Behav. 14, 226–235. (doi:10.1016/0003-3472(66)80076-3)

34. Dunn PK. 2017 Tweedie: evaluation of Tweedie exponential family models. R package version 2.3.

35. Zeliea A, Kleiber C, Jackman S. 2008 Regression models for count data in R. J. Stat. Softw. 27, 1–25.

36. R Development Core Team. 2020 R: a language and environment for statistical computing. Vienna, Austria: R Foundation for Statistical Computing. See http://www.R-project.org/.

37. Maia R, Eliason CM, Bitton P-P, Doucet SM, Shawkey MD. 2013 pavo: An R package for the analysis, visualization and organization of spectral data. Methods Ecol. Evol. 4, 609–613. (doi:10.1111/2041-210X.12069)

38. Neville MC et al. 2014 Male-specific fruitless isoforms target neurodevelopmental genes to specify a sexually dimorphic nervous system. Curr. Biol. 24, 229–241. (doi:10.1016/j.cub.2013.11.035)

39. Robertson KA, Monteiro A. 2005 Female Bicyclus anynana butterflies choose males on the basis of their dorsal UV-reflective eyepatch pupils. Proc. R. Soc. B 272, 1541–1546. (doi:10.1098/rspb.2005.3142)

40. Huq M, Bhawardj S, Monteiro A. 2019 Male Bicyclus anynana butterflies choose females on the basis of their ventral UV-reflective eyepatch centers. J. Insect. Sci. 19, 25. (doi:10.1093/jisesa/izx014)

41. Ringo JM. 2002 Hormonal regulation of sexual behavior in insects. In Hormones, brain and behavior (eds DW Pfaff, M Joels), pp. 93–114. London, UK: Academic Press.

42. Barth RH, Lester LJ. 1973 Neuro-hormonal control of sexual behavior of decapitated females. In Insect. Biochem. Mol. Biol. 19, 609–617. (doi:10.1016/0020-708X(73)90048-3)

43. Ishimoto H, Sakai T, Kitamoto T. 2009 Ecdysone signaling regulates the formation of long-term courtship memory in adult Drosophila melanogaster. Proc. Natl Acad. Sci. USA 106, 6381–6386. (doi:10.1073/pnas.0810213106)

44. Westerman EL, Drucker CB, Monteiro A. 2014 Male and female mating behavior is dependent on social context in the butterfly Bicyclus anynana. J. Insect Behav. 27, 478–495. (doi:10.1007/s10590-014-9441-9)

45. Barek H, Venkasa A, Sugumaran M. 2018 Drosophila melanogaster has the enzymatic machinery to make the melanin component of neuromelanin. Pigment Cell and Melanoma Res. 31, 683–692. (doi:10.1111/pcm.12709)

46. Yamamoto S, Seto ES. 2014 Dopamine dynamics and signaling in Drosophila: an overview of genes, drugs and behavioral paradigms. Pigment Cell and Melanoma Res. 63, 107–119. (doi:10.1538/pcmr.63.107)

47. Zhang SX, Rogulja D, Crookmore MA. 2016 Dopaminergic circuitry underlying mating drive. Neuron 91, 168–181. (doi:10.1016/-neuron.2016.05.020)

48. Xu X et al. 2011 Structure and function of a ‘yellow’ protein from saliva of the sand fly Lutzomyia longipalpis that confers protective immunity against Leishmania major infection. J. Biol. Chem. 286, 32 383–32 393. (doi:10.1074/jbc.m111.269904)

49. Verlinden H. 2018 Dopamine signaling in locusts and other insects. Insect. Biochem. Mol. Biol. 97, 40–52. (doi:10.1016/j.ibmb.2018.04.005)

50. Rauschenbach IY, Bogomolova EV, Karpova EK, Adonyeva NV, Faddeeva NV, Mueshanov PN, Grunenko NE. 2011 Mechanisms of age-specific regulation of dopamine metabolism by juvenile hormone and 20-hydroxyecdysone in Drosophila females. J. Comp. Physiol. B 181, 19–26. (doi:10.1007/s00360-010-0512-8)

51. Zilberman-Peled B, Ren B, Gross A, Finberg JP, Gottlief Y. 2006 A possible new role for fish retinal serotonin-N-acetyltransferase-1 (AANAT1): dopamine metabolism. Brain Res. 1073–1074, 220–228. (doi:10.1016/j.brainres.2005.12.028)

52. Rauschenbach IY, Lautkina OV, Alekseev AA, Adonyeva NV, Bogomolova EV, Grunenko NE. 2012 Dopamine effect on 20-hydroxyecdysone level is mediated by juvenile hormone in Drosophila females. Dokl. Biochem. Biophys. 446, 263–265. (doi:10.1134/S1607672912050171)

53. Ellis LL, Carney GE. 2010 Mating alters gene expression patterns in Drosophila melanogaster male heads. BMC Genom. 11, 558. (doi:10.1186/1471-2164-11-558)

54. Grunenko NE, Karpova EA, Alekseev AA, Chentsova NA, Shumnya LV, Ushakova MA, Rauschenbakh IY. 2006 An increase in the dopamine level accelerates sexual maturation of Drosophila melanogaster deficient in the juvenile hormone. Dokl. Biol. Sci. 406, 88–90. (doi:10.1134/s106451720601025x)

55. Harano K, Sasaki K, Nagao T, Sasaki M. 2008 Influence of age and juvenile hormone on brain dopamine level in male honeybee (Apis mellifera): association with reproductive maturation. J. Insect. Physiol. 54, 848–853. (doi:10.1016/j.jiphysiol.2008.03.003)

56. Grunenko NE, Karpova EA, Chentsova NA, Adonyeva NV, Rauschenbach IY. 2009 20-Hydroxyecdysone and juvenile hormone influence tyrosine hydroxylase activity in Drosophila females under normal and heat stress conditions. Arch. Insect. Biochem. Physiol. 72, 263–272. (doi:10.1002/arch.20337)

57. Connahs H, Tan EJ, Ter YT, Dion E, Matsuoka Y, Leavey B, Bear A, Monteiro A. 2022 Data from: The yellow gene regulates behavioural plasticity by repressing male courtship in Bicyclus anynana butterflies. Dryad Digital Repository. (doi:10.5061/dryad.9np3shq94)

58. Connahs H, Tan EJ, Ter YT, Dion E, Matsuoka Y, Bear A, Monteiro A. 2022 The yellow gene regulates behavioural plasticity by repressing male courtship in Bicyclus anynana butterflies. Figshare.