Neff & Svensson [1] recently reviewed the literature relating to the theoretical modelling of alternative mating tactics. Their review presents insightful future perspectives on how genomics could unveil the molecular genetics and proximate mechanisms of tactic expression, and we fully agree with their statements that a better understanding of the field can be achieved through the integration between quantitative and molecular genetics. However, we feel that it is important to draw attention to two significant oversights: their review (i) neglects previous studies that provided theoretical and empirical contributions to the understanding of the genetics underlying the conditional expression of alternative mating tactics (i.e. conditional strategies) and (ii) presents ideas that have been previously published (by other authors) as a new ‘unified theory for the evolution and phenotypic expression of alternative mating tactics’. Below we elaborate why we believe that Neff and Svensson’s verbal model (‘the conditional alternative strategy’) is not a new model, but is instead based on a misinformed view that current models do not account for genetic variation underlying conditional strategies. As a result, Neff and Svensson’s ‘conditional alternative strategy’ unnecessarily muddles our understanding of conditional strategies.

Phenotypic plasticity is a ubiquitous evolutionary phenomenon [2] and nowhere perhaps, is plasticity more apparent than when single genotypes can produce different alternative phenotypes depending upon environmental conditions [3]. Such polyphenisms are synonymous with conditional strategies and have been described in a wide range of traits, including the presence, shape and colour of morphological traits, alternative mating tactics, diet, sex and caste determination, as well as paedomogenesis and diapause [4]. Understanding their genetic architecture and evolution is therefore vital for our understanding of these important adaptations. Dawkins’ conditional evolutionary stable strategy model [5] provided an important theoretical framework for understanding polyphenisms, but also conveyed the message that genetic polymorphism is not necessary for a conditional strategy to be evolutionary stable. This message was an attempt to distinguish conditional strategies from ‘alternative strategies’, which represent a genetic polymorphism with Mendelian inheritance. However, it also led numerous authors to erroneously associate conditional strategies with complete genetic monomorphism [1,6,7]. It is well established, however, that underlying genetic variation is prevalent in plastic traits [8,9], and its role in polyphenisms has been modelled [10,11] and supported empirically numerous times [12–15]. The conditional strategy was recently criticized based on this false assumption of genetic monomorphism [7]—a misconception that was subsequently laid to rest [16]. Neff and Svensson do cite this [16] correction of the record, but in our view failed to convey an understanding of its content, because the assumption of monomorphism re-appears as the driver of Neff & Svensson’s [1] ‘new model’.

Sewall Wright [17,18] pioneered the idea that discontinuous traits could result from continuous polygenic variation, coupled with a threshold mechanism that generates discontinuity in trait expression. Such traits were named
threshold traits [19], and the concept of continuous polygenic variation underlying their expression became the central idea of the ‘liability model’ of quantitative genetics [20,21]. This model assumes liability is a continuously variable quantitative trait, influenced by genetic and environmental factors. Individuals produce one or the other of two alternative phenotypes depending on whether their liability exceeds some fixed threshold [20]. Wright’s model can be modified to account for the alternative tactics of conditional strategies by allowing the position of the threshold relative to the continuous distribution of liabilities to be a function of the environmental cues that influence the expression of the alternative tactics. Thus, according to this ‘environmental threshold model’, additive genetic variation, not genetic monomorphism, underlies the expression of the alternative tactics of the conditional strategy [4,10,16].

Unarguably, the view that genetic variation can influence the conditional expression of alternative tactics is neither new nor commonly disregarded—the three main references on the quantitative genetics of conditional strategies, namely Hazel et al. [10], Roff [4] and Hazel et al. [11] sum more than 360 citations in Web of Science to date. Despite briefly citing some of these studies, Neff & Svensson [1] disregard the ideas and models in them, and then present the conceptualization of genetic variation for switch points as if it were a new idea (see their section ‘Genetic and plastic alternative mating tactics’). Not only has this idea been described previously, it has also been recently reviewed [14,16] and empirically supported [12,13,22–25]. Interestingly, Neff & Svensson [1] acknowledge and cite previous work that empirically supported [12]. Unfortunately, most of these studies were also overlooked by Neff & Svensson [1].

In conclusion, the game-theoretical models of conditional and alternative strategies do not need to be fused in order to accommodate genetic variation for conditional polymorphisms, because this has already been comprehensively done under the umbrella of the environmental threshold model for conditional strategies. Presenting this fusion and labelling it as a putative new unified theory [1] is unnecessary and mathematically unsupported. Much progress has been done in the last few decades on the genetic basis of conditional strategies, and it is clear that genetic variation in conditional strategies is prevalent in natural populations. The genetic monomorphism-oriented criticism of conditional strategies present in Neff & Svensson [1] (see also [7]) adds nothing to the progress already made [10,11,14,16]. Finally, since a new verbal model is proposed to do what previous mathematically supported models have already been doing for years, we believe that the ‘conditional alternative strategies’ model is unjustified, and even muddies the water when it unnecessarily fuses alternative strategies with conditional strategies.

References


18. Wright S. 1934 The results of crosses between inbred strains of guinea pigs, differing in number of digits. *Genetics* 19, 537 – 551.


