A global meta-analysis reveals higher life-history phenotypic

variation in urban birds than in their non-urban neighbours

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Author contributions

PC-L, MJT, DR, AC and DMD conceived the study. PC-L, AS-T, CJB and DMD performed

the literature search and PC-L extracted effect sizes from suitable published papers. PC-L

and YH performed all statistical analysis with advice from AS-T. PC-L wrote the first draft of

the manuscript with input from MJT, AS-T, DR, AC and DMD. All authors read and revised

the manuscript.

Data availability

All R scripts and datasets needed to reproduce the analyses presented in this paper are

available at: https://github.com/PabloCapilla/meta-analysis_variation_urban.

Abstract

Cities pose a major ecological challenge for wildlife worldwide. Phenotypic variation, which can result from underlying genetic variation or plasticity, is an important metric to understand eco-evolutionary responses to environmental change. Recent work suggests that urban populations might have higher levels of phenotypic variation than non-urban counterparts. This prediction, however, has never been tested across taxa nor over a broad geographical range. Here, we conduct a meta-analysis of the avian literature to compare urban *versus* non-urban means and variances in breeding phenology (i.e., lay date) and reproductive effort (i.e., clutch size, number of fledglings). First, we show that urban populations reproduce earlier and have smaller broods than non-urban conspecifics. Second, we show that urban bird populations have higher phenotypic variation in laying date than non-urban populations. This result arises from differences in phenological variation between populations within breeding seasons, conceivably due to higher landscape heterogeneity in urban habitats. Higher variation in reproductive timing in birds subject to anthropogenic disturbance might be a consequence of plastic responses or higher genetic variation in phenology in urban populations. Our results reveal a novel effect of urbanisation on animal life-history traits with important implications for our understanding of how species adapt to urban environments.

Introduction

Humans have drastically changed environmental conditions on Earth (Hendry *et al.* 2017). The footprint of human activity is most pronounced in urban environments, where microclimatic conditions, biogeochemical cycles and sensory landscapes are considerably different from those in non-urban environments (Grimm *et al.* 2008). Perhaps not surprisingly, multiple shifts in animal and plant phenotypes have been associated with the novel conditions and selective pressures found in cities (Johnson & Munshi-South 2017; Lambert *et al.* 2020). Indeed, numerous studies have reported divergent phenotypes between urban and non-urban populations across the globe (e.g., Alberti *et al.* 2017; Diamond *et al.* 2018; Campbell-Staton *et al.* 2020; reviewed in Johnson & Munshi-South 2017; Lambert *et al.* 2020; Diamond & Martin 2021). The extent to which populations can adapt to urban environments could be partly associated with how urbanisation affects their phenotypic variation (Thompson *et al.* 2022). Phenotypic variation is tightly linked to eco-evolutionary processes: it is an essential condition for current selection, it results from past selection pressures, and it depends on gene flow and phenotypic plasticity. As such, assessing how urbanisation affects phenotypic variation can help us understand the potential for future phenotypic changes in urban environments and the eco-evolutionary implications of such changes (Thompson *et al.* 2022).

Urban evolutionary ecologists have traditionally focused on differences in mean trait values between urban and non-urban populations, typically finding altered phenology, morphology, behaviour and reproductive performance in urban populations (or populations exposed to urban-like factors) compared to non-urban counterparts (Chamberlain *et al.* 2009; Vaugoyeau *et al.* 2016; Sepp *et al.* 2018; Putman & Tippie 2020; Sanders *et al.* 2021). However, no study to date has explicitly investigated how urbanisation affects phenotypic variation in reproductive traits at a global scale.

Recent single-species studies suggest that phenotypic variation could be affected by

urbanisation (Caizergues *et al.* 2018; Gorton *et al.* 2018; Thompson *et al.* 2022). For

example, in species with limited dispersal ability (i.e., whose dispersal occurs at a smaller

scale than the scale at which the urban habitat varies), adaptation to local conditions could

increase phenotypic variation within the urban matrix. Findings from urban and non-urban

meta-populations of the common ragweed (*Ambrosia artemisiifolia*) are consistent with this

prediction as inter-population variation in several fitness proxies was greater in urban

environments compared to non-urban environments (Gorton *et al.* 2018). A meta-analysis of

selection strength found weaker selection occurring in human-disturbed populations (Fugère

& Hendry 2018; although this study did not specifically test the effect of urbanisation on

selection strength and only included one study directly associated with urbanisation), which if

extrapolated to the urban context, could lead to higher phenotypic variation in urban

populations compared to their non-urban counterparts. Overall, these studies converge with

the notion that urban populations could display higher levels of phenotypic variation, due to

several eco-evolutionary processes reviewed in (Thompson *et al.* 2022). These findings also

highlight that the extent to which urbanisation may impact phenotypic variation likely

depends on the interplay between the temporal and spatial scale at which environmental

conditions fluctuate in the urban habitat, as well as on the species' longevity and dispersal

ability (Thompson *et al.* 2022).

The temporal scale at which differences in phenotypic variation between urban and non-

urban habitats manifest can help us evaluate their ecological causes, and is likely to

determine the eco-evolutionary implications of increased phenotypic variation in urban

habitats (Thompson *et al.* 2022). First, urban populations could display higher phenotypic

variation than non-urban populations within a given breeding season (as a result, for

example, of consistent differences in landscape heterogeneity between habitats; Pickett *et*

al. 2017). Second, urban populations could display higher phenotypic variation than non-

urban populations due to larger yearly fluctuations in environmental conditions (if, for

example, urban populations are less resilient to changes in weather conditions), with or

without differences in phenotypic variation between urban and non-urban populations within

a given season. In the latter scenario, similar levels of phenotypic variation would be

exposed to natural selection in short-lived species (e.g., annual species).

Urban environments have been referred to as spatially more heterogeneous than non-urban

habitats of the same geographical area (Pickett *et al.* 2017). Urban habitat heterogeneity

could increase phenotypic variation compared to adjacent non-urban habitats if, for example,

urban organisms change their phenotype according to local environmental conditions (e.g.,

through both developmental and later-life phenotypic plasticity). The empirical assessment of

this idea, however, largely depends on the scale at which urban habitat heterogeneity is

measured, the spatial scale at which the organism of interest operates and the heterogeneity

of the non-urban habitat of reference (Pickett *et al.* 2017; Uchida *et al.* 2021). For example, a

megacity may be spatially heterogeneous, containing a diverse array of habitats (e.g.,

multiple urban parks with different ecological conditions, a varying level of impervious

surface, etc.), and, thus, be overall vastly more heterogeneous than a neighboring non-

urban habitat. However, a given species of interest could reduce the range of environmental

conditions that it experiences through matching habitat choice (e.g., (Muñoz *et al.* 2014)),

limiting the potential effect of urban habitat heterogeneity on phenotypic variation. Therefore,

measuring habitat heterogeneity at different spatial scales will be paramount to understand

the potential association between habitat heterogeneity and increased phenotypic variation

in urban areas.

Here, we investigate how urbanisation impacts mean phenotypic values and phenotypic

variation using a meta-analysis of 409 paired urban and non-urban comparisons of avian

life-history traits (laying date, clutch size and number of fledglings) published between 1958

and 2020 including 33 bird species (Figure 1). We use within species urban – non-urban

comparisons to investigate the following questions: i) Is urbanisation associated with shifts in

mean life-history traits? ii) Is urbanisation associated with changes in variation in life-history traits? iii) What is the temporal and spatial scale at which urbanisation correlates with changes in phenotypic variation? Based on previous research (Chamberlain *et al.* 2009; Sepp *et al.* 2018), we predict that urban bird populations display on average earlier phenology, smaller clutch size and lower number of fledglings than non-urban populations. We also predict increased phenotypic variation in urban populations compared to non-urban populations for all three traits examined (see above; and reviewed in Thompson *et al.* 2022). Lastly, we disentangle urban effects on phenotypic variation across different temporal and spatial scales, suggesting an ecological mechanism for the effects of urbanisation on avian phenotypic variation. This study provides, for the first time, meta-analytical evidence that urban conditions can magnify phenotypic variation in avian life-history traits and highlights the potential role of increased habitat heterogeneity in urban areas as an ecological mechanism underlying this effect.

Material and methods

Literature review

We began our literature search by inspecting two published reviews on the impact of urbanisation on avian biology (Chamberlain *et al.* 2009; Sepp *et al.* 2018). As we were mainly interested in how phenology and reproduction were affected by urbanisation, we identified papers cited in Chamberlain *et al.* (2009) (N = 37) and Sepp *et al.* (2018) (N = 32) that could contain either data, or mean and variance estimates for first clutch laying initiation (hereafter laying date), clutch size and number of nestlings fledged per breeding attempt (hereafter number of fledglings), for paired urban and non-urban populations. Then, we performed four searches of the Web of Science Core Collection on the 27th of October 2020 (databases covered: SCI-EXPANDED – 1900-present, SSCI – 1956-present, A&HCI – 1975-present, BKCI-S – 2005-present, BKCI-SSH – 2005-present and ESCI – 2015-present) to recover studies published since 1900 and including all languages and all document types.

We performed the following four searches: (**1**) TS=("urban*" AND ("bird*" OR "aves" OR "avian" OR "ornithol*" OR "passerine*" OR "passeriform*" OR "songbird*" OR *list of bird genera*) AND ("laying date" OR "lay date" OR "first egg" OR "clutch size" OR "eggs laid" OR "number of eggs" OR "fledgling*" OR "fledging" OR "reproductive success" OR "fitness")); (**2**) TS=("urban*" AND "bird" AND "laying date"); (**3**) TS=("urban*" AND "bird" AND "clutch size"); (**4**) TS=("urban*" AND "bird" AND "fledglings"). The *list of avian genera* in the first search string consisted of a list of all avian genera and can be found in Supplementary text (see also acknowledgements). Searches 2-4 were contained within search 1; however, they were not restricted to avian genera, hence potentially capturing additional references that did not mention the scientific name of the study species. These searches found 892, 71, 198 and 167 studies, respectively, which we combined with the studies identified from Chamberlain *et al.* (2009) and Sepp *et al.* (2018) to create a list of 1,397 studies (Figure S1). We then de-duplicated this list using the R package 'revtools' (using exact matching of study titles in function '*find_duplicates*', v0.4.1; Westgate 2019) and by manually inspecting all study titles and author lists. Our final list contained 940 unique studies (Figure S1). We screened each study by reading their title and abstract. If the title and/or abstract indicated that the paper could fit our requirements for data collection (see below), we read the study fully, aiming to extract mean, standard deviation (SD) and sample size (n) of our life-history traits of interest for urban and non-urban bird populations. If SD was not available but authors provided SE, the former was calculated as: $SD = SE \times \sqrt{n}$. Mean and SD were extracted from data quartiles and medians in four effect sizes from two studies following Luo *et al.* (2016) and Shi *et al.* (2020). When available, we extracted estimates per breeding season (i.e., papers often reported mean, SD and n for urban and non-urban populations in multiple breeding seasons). If a study reported incomplete information for inclusion in our meta-analysis (e.g., mean was provided but not SD or SE), we contacted the authors to ask for this missing information (a complete list of authors that provided estimates can be found in the acknowledgments).

Criteria for inclusion

We were interested in investigating the effects of urbanisation on life-history traits, with a particular interest in testing the association between urbanisation and, mean and variation in trait values. Paired urban – non-urban designs, where an urban population is compared to a non-urban adjacent population, are a powerful approach to identify the effects of urban living while controlling for temporal and geographical variation, as well as large-scale genetic structure among populations. Therefore, we included studies if they (i) compared geographically close (i.e., paired) urban and non-urban populations and (ii) reported laying date of the first clutches of the season, clutch size or number of nestlings for the same breeding season across both habitats. When multiple populations were compared along a gradient of urbanisation, we extracted estimates for the two populations at the extremes of the gradient (i.e., most urban and least urban populations). When studies combined estimates across several breeding seasons, we included them in our meta-analysis if urban and non-urban populations had been sampled in the same breeding seasons.

Initially, our data set contained 427 paired urban – non-urban estimates from 33 bird species and 65 studies. Of these, three observations were removed due to missing sample sizes, and 11 observations were removed because their sample size was one (which precludes the calculation of a mean and their SD). Four observations were removed because they reported a SD of zero (these indeed had very low sample sizes: 3, 2, 7, 2 data points). One effect size for laying date, nine effect sizes for clutch size and 13 effect sizes for number of fledglings from five studies contained missing SD data. Mean and SD are often strongly correlated (Nakagawa & Schielzeth 2013), including in our dataset (Figure S2). Therefore, we imputed these 23 missing SD values based on linear regressions of log mean against log standard deviation (not including imputed values in our meta-analysis produced quantitatively very similar results; k = 386 comparisons not including imputed SD values). Our final data set included 409 comparisons between paired urban and non-urban populations from 33 bird species and 65 studies (Figure 1; refs.: Perrins 1965; Middleton 1979; Schmidt & Steinbach

1983; Dhondt *et al.* 1984; Eden 1985; Stout *et al.* 1998; Boal & Mannan 1999; Mcgowan 2001; Schoech & Bowman 2001; Solonen 2001; Antonov & Atanasova 2003; Rollinson & Jones 2003; Liven-Schulman *et al.* 2004; Millsap *et al.* 2004; Conway *et al.* 2006; Beck & Heinsohn 2006; Mennechez & Clergeau 2006; Isaksson & Andersson 2007; Kelleher & O'Halloran 2007; Pande *et al.* 2007; Schoech *et al.* 2007; Hinsley *et al.* 2008; Isaksson *et al.* 2008; Newhouse *et al.* 2008; Solonen & Ursin 2008; Ibáñez-Álamo & Soler 2010; Berardelli *et al.* 2010; Shustack & Rodewald 2011; Seress *et al.* 2012, 2018, 2020; Stracey & Robinson 2012; Cardilini *et al.* 2013; Brahmia *et al.* 2013; Morrissey *et al.* 2014; Solonen & Hildén 2014; Sumasgutner *et al.* 2014; Gahbauer *et al.* 2015; Glądalski *et al.* 2015, 2016a, b, 2017; Lin *et al.* 2015; Wawyrzyniak *et al.* 2015; Bailly *et al.* 2016; Minias 2016; Perlut *et al.* 2016; Lee *et al.* 2017; Meyrier *et al.* 2017; Biard *et al.* 2017; Pollock *et al.* 2017; Preiszner *et al.* 2017; Thornton *et al.* 2017; Capilla-Lasheras *et al.* 2017; Gryz & Krauze-Gryz 2018; Bobek *et al.* 2018; Caizergues *et al.* 2018; de Satgé *et al.* 2019; Hajdasz *et al.* 2019; Kettel *et al.* 2019; Rosenfield *et al.* 2019; Welch-Acosta *et al.* 2019; Jarrett *et al.* 2020; Luna *et al.* 2020; Partecke *et al.* 2020). Of these 409 comparisons, 151 corresponded to comparisons of (first clutch) laying date (N = 32 studies), 124 corresponded to comparisons of clutch size (N = 42 studies) and 134 were comparisons of number of fledglings (N = 44 studies) (Figure S3). Last, there were a total of 376 comparisons for single years (N = 47 studies) and an additional 33 comparisons included estimates across multiple years of study (N = 18 studies).

Meta-analytic effect sizes

Laying date data was standardised across studies by coding it as days after January 1. In every case, mean laying date estimates across habitats fell within the same calendar year. For each of these three life-history traits, we computed the log response ratio (lnRR) and the log coefficient of variation ratio (lnCVR) to investigate differences in mean values and variability between urban and non-urban populations (Hedges *et al.* 1999; Nakagawa *et al.* 2015; Senior *et al.* 2020). We calculated lnRR and lnCVR along with their associated

sampling variances (Nakagawa *et al.* 2015) using the R function '*escalc*' in the 'metafor' R

package (v2.5-57; Viechtbauer 2010). Both lnRR and lnCVR were calculated in a way that

positive values meant higher estimates in urban populations compared to their non-urban

counterparts. Urban and non-urban populations can differ in mean and variance values at

the same time (e.g., Taylor's Law; Nakagawa & Schielzeth 2013; Cohen & Xu 2015), as was

observed in our data (Figure S2). Therefore, we chose lnCVR over lnVR (i.e., log total

variation ratio; Nakagawa *et al.* 2015) as the former is designed to account for the mean–

variance relationship (Nakagawa *et al.* 2015; Senior *et al.* 2020). However, we also show

results for the log total variation ratio in Figure S5.

Quantifying habitat heterogeneity and urban index

We calculated habitat heterogeneity from the 3CS LC (Copernicus Climate Change Service

Land Cover) and the ESA-CCI LC (European Space Agency-Climate Change Initiative Land

Cover) land cover products ("ESA. Land Cover CCI Product User Guide" 2017; "ESA. 3CS

Land Cover Product User Guide" 2020). These datasets provide methodologically consistent

land cover per year and gridded maps from 1992 to 2019, with a global coverage and a

spatial resolution of circa 300 m per pixel (0.002778° or 10 arcseconds). Each pixel in the

products is classified as one of the 22 land cover categories defined by the UN-FAO-LCCS

(United Nations Food and Agriculture Organization Land Cover Classification System)

("ESA. Land Cover CCI Product User Guide" 2017; "ESA. 3CS Land Cover Product User

Guide" 2020). From those studies included in our main meta-analysis, we extracted the

coordinates of their urban and non-urban populations (N = 30 studies with coordinates

provided for urban and non-urban populations). Then, we sampled the landscape of every

study by extracting the number of pixels belonging to each land cover category around each

urban and non-urban location (i.e., within a circular buffer around each location). The

extraction was performed for several buffer radii from 250 m to 5000 m in intervals of 250 m.

Landscape heterogeneity was calculated as the effective number of land covers present in

each buffer and computed as the exponential of the Shannon-Wiener diversity index (i.e.,

Hill's numbers for $q = 1$) (Hill 1973; Chao *et al.* 2014), resulting into a measure that not only

accounts for the absolute richness of land cover categories but also weights in the relative

abundance of each category. Urban index was calculated as the proportion of each buffer

area categorized as an 'urban' land cover type. Land cover data were processed and

analysed using R (v.4.0.3; R Core Team 2021). Geospatial vectorial operations were

conducted utilising the 'sf' R package (v.1.0-7; Pebesma 2018) while raster extraction were

performed with the 'raster' R package (v.3.5-15; Hijmans 2020). All geospatial analyses were

performed in the WSG 1984 projected Coordinate Reference Systems, EPSG: 6326. We

could retrieve location information for 250 urban *versus* non-urban comparisons for laying

date, clutch size and number of fledglings, from 12 species ad 30 studies between 1992 and

2017 (land cover data were not available before 1992; see above).

Meta-analyses and random effects

We handled the data sets, ran all analyses and produced visualisations using R (v.4.0.3; R

Core Team 2021). To evaluate the overall effect of urbanisation on avian life-history traits,

we fitted phylogenetic multilevel (intercept-only) meta-analyses for each response term (i.e.,

lnRR and lnCVR) combining the three life-history traits (i.e., laying date, clutch size and

number of fledglings). Both meta-analytic models estimated four random intercept effects,

publication identity (i.e., among-study variation), population identity, phylogeny (more details

below) and species identity (i.e., among-species variation not explained by phylogeny), as

well as a residual variance term. We estimated total heterogeneity (I^2) and the proportion of

variation explained by each random term in our meta-analytic (intercept-only) models

following (Nakagawa & Santos 2012; Senior *et al.* 2016) and as implemented in the R

function '*i2_ml'* ('orchaRd' R package v.0.0.0.9000; Nakagawa *et al.* 2020).

Phylogenies

Phylogenetic trees were extracted from the Open Tree of Life (Hinchliff *et al.* 2015; Rees &

Cranston 2017), using the interface provided by the R package 'rotl' (v3.0.10; Michonneau et

al. 2016, OpenTreeOfLife et al. 2019). We calculated tree branch length (Grafen 1989) and

generated a phylogenetic correlation matrix to include in our phylogenetic meta-analytic

models (Figure 1). We assessed the phylogenetic signal in our meta-analysis based on the

proportion of variation explained by the phylogeny (i.e., $I^2_{\text{phylogeny}}$; e.g., Cinar *et al.* 2022).

Modelling heterogeneous variances and correlations among traits

Laying date, clutch size and number of fledglings are often correlated in bird species (Rowe

et al. 1994; Dunn & Møller 2014). To assess whether urbanisation is associated with

correlated responses across life-history traits and the robustness of our results to the

existence of these correlations, we built trivariate meta-analytic models of lnRR and lnCVR

that allowed us to simultaneously estimate trait-specific means (i.e., one intercept for each

trait – Equation 1), trait-specific residual variances (i.e., one residual variance for each trait –

Equation 1 & Equation 2) and trait-specific among-study variances and correlation among

traits (Equation 1 & Equation 3). Including the random-effect terms detailed above, our

model with heterogeneous variances and among-study correlations among traits can be

written as: (we have omitted the term associated with sampling variance for simplicity – see

Nakagawa et al. (2015) for more details)

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$$y_i = \begin{bmatrix} \mu_{LD} \\ \mu_{CS} \\ \mu_{NF} \end{bmatrix} + \begin{bmatrix} \varepsilon_{i-LD} \\ \varepsilon_{i-CS} \\ \varepsilon_{i-NF} \end{bmatrix} + \begin{bmatrix} \tau_{t-LD} \\ \tau_{t-CS} \\ \tau_{t-NF} \end{bmatrix} + v_y + a_l + h_w \text{ (Equation 1)}$$

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$$\begin{matrix} \varepsilon_{i-LD} \\ \varepsilon_{i-CS} \\ \varepsilon_{i-NF} \end{matrix} \sim N(\begin{bmatrix} 0 \\ 0 \\ 0 \end{bmatrix}, \begin{bmatrix} \sigma^2_{\varepsilon-LD} \\ \sigma^2_{\varepsilon-CS} \\ \sigma^2_{\varepsilon-NF} \end{bmatrix}) \text{ (Equation 2)}$$

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$$\begin{matrix} \tau_{t-LD} \\ \tau_{t-CS} \\ \tau_{t-NF} \end{matrix} \sim N(\begin{bmatrix} 0 \\ 0 \\ 0 \end{bmatrix}, \begin{bmatrix} \sigma^2_{t-LD} & p_{LD-CS}\,\sigma_{t-LD}\,\sigma_{t-CS} & p_{LD-NF}\,\sigma_{t-LD}\,\sigma_{t-NF} \\ & \sigma^2_{t-CS} & p_{CS-NF}\,\sigma_{t-CS}\,\sigma_{t-NF} \\ & & \sigma^2_{t-NF} \end{bmatrix}) \text{ (Equation 3)}$$

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$$v_y \sim N(0, \sigma^2_v) \text{ (Equation 4)}$$

$$a_l \sim N(0, A\sigma_a^2) \text{ (Equation 5)}$$

$$h_w \sim N(0, \sigma_h^2) \text{ (Equation 6)}$$

Where y_i is the statistic of interest (lnRR or lnCVR) for the ith effect size ($i = 1,2,3, \dots, k$; where k is the number of the effect sizes included in the analysis - i.e., number of urban – non-urban paired comparisons). 'LD', 'CS', 'NF' refer to overall means (μ), variances (σ) and correlations (ρ) involving effect sizes for laying date ('LD'), clutch size ('CS') and number of fledglings ('NF'). ε_i is the residual deviation for the ith observation, which is assumed to follow a normal distribution with mean 0 and variance $\sigma_{\varepsilon-LD}^2$, $\sigma_{\varepsilon-CS}^2$, $\sigma_{\varepsilon-NF}^2$ for laying date, clutch size and number of fledglings, respectively. τ_{t-LD}, τ_{t-CS}, and τ_{t-NF} are the deviations from the mean associated with the tth study and trait ('LD', 'CS', or 'NF'), each following a multivariate normal distribution with mean of 0 and variance-covariance structure detailed in Equation 5 (p provides the correlations between τ_{t-LD}, τ_{t-CS}, and τ_{t-NF}). v_y provides the deviation from the overall mean associated with the yth population. a_l is the phylogenetic effect for the lth species, which follows a normal distribution with mean equal to 0 and variance-covariance structure given by σ_a^2, the variance of the phylogenetic effect, and A, a l by l matrix of distances between species calculated from a phylogenetic tree (details above). h_w captures among species variation not explained by the phylogenetic effect and follows a normal distribution around 0 and variance σ_h^2.

We compared models with different constraints in the parameters of the variance-covariance structure in Equation 3 to assess the strength of evidence for heterogeneous variances and correlations among traits. We fitted these trivariate meta-analytic models in the 'metafor' R package (v2.5-57; Viechtbauer 2010) using maximum likelihood and compared model AIC (Akaike information criterion; Burnham *et al.* 2011). We then calculated a ΔAIC value for

each model (i.e., the difference in AIC between a given model and the model with lowest

AIC) and used this value to assess the strength of evidence for a given variance-covariance

structure. We fitted models with the following constraints in the variance-covariance

structure:

(i)   Single variance across traits and zero covariances:

$\sigma^2_{t-LD} = \sigma^2_{t-CS} = \sigma^2_{t-NF}$; and all $p = 0$

(ii)   Compound symmetric variance-covariance matrix:

$\sigma^2_{t-LD} = \sigma^2_{t-CS} = \sigma^2_{t-NF}$; and $p_{LD-CS} = p_{LD-NF} = p_{CS-NF}$

(iii)   Heteroscedastic compound symmetric variance-covariance matrix:

$\sigma^2_{t-LD}, \sigma^2_{t-CS}$ and σ^2_{t-NF} can vary freely but $p_{LD-CS} = p_{LD-NF} = p_{CS-NF}$

(iv)   Diagonal variance-covariance matrix:

$\sigma^2_{t-LD}, \sigma^2_{t-CS}$ and σ^2_{t-NF} can vary freely but all $p = 0$

(v)   Unstructured variance-covariance matrix

$\sigma^2_{t-LD}, \sigma^2_{t-CS}$ and σ^2_{t-NF} can vary freely, as well as p_{LD-CS}, p_{LD-NF} and p_{CS-NF} (i.e.,

correlation among traits).

Within- and between-breeding season differences in phenology and life-history traits

between urban and non-urban population

Urban and non-urban populations may differ in both within- and between-breeding season

variation in phenology and life-history traits. However, differences in variation for these two

temporal scales are likely generated by contrasting ecological and evolutionary processes

(Thompson *et al.* 2022). To disentangle processes operating at these two temporal scales,

we performed additional meta-analyses including 1) urban – non-urban comparisons within

breeding seasons (k = 376 comparisons in the original data set) and 2) urban – non-urban

comparisons between breeding seasons (i.e., combining all within-breeding season

estimates from a given study; k = 33 comparisons present in the original data set, plus 68

additional comparison calculated from within-breeding season estimates; see below). When

a given study reported estimates across multiple breeding seasons, we calculated between-

breeding season mean and variance as:

$$\bar{x}_{among-season} = \sum_{i=1}^{g} \bar{x}_i \frac{n_i}{N}; where, N = \sum_{i=1}^{g} n_i \text{ (Equation 10)}$$

$$S_{among-season}^2 = \sum_{i=1}^{g} n_i s_i^2 + \sum_{i=1}^{g} n_i \left(\bar{x}_i - \bar{x}_{among-season} \right)^2 \text{ (Equation 11)}$$

Where, $\bar{x}_{among-season}$ and $S_{among-season}^2$ are mean and variance across multiple breeding

seasons. g is the total number of breeding seasons reported by a given study; $\bar{x}_i$, s_i^2, n_i, are

mean, variance and sample size for each breeding season i. $\bar{x}_{among-season}$ for a given study

is, therefore, the weighted average across breeding seasons (Equation 10); whereas

$S_{among-season}^2$ for a given study is the weighted sum of within-season variances (first term in

equation 11) and between-season mean variances (second term in Equation 11).

Assessing the effect of urbanisation and habitat heterogeneity on differences in

phenotypic variation between habitats

Using the subset of studies which allowed quantification of urban index in urban and non-

urban populations (see above), we first verified that urban index was indeed higher for urban

populations than for non-urban populations. We compared urban index in urban and non-

urban populations at different spatial scales via linear models. Then, to assess whether the

increase in phenotypic variation in urban habitats was predicted by habitat heterogeneity

and/or urban index, we ran an additional meta-regression to explain differences in

phenotypic variation between urban and non-urban populations (i.e., lnCVR), where the

difference in habitat heterogeneity and urban index between urban and non-urban

populations were included as continuous moderators. This meta-regression included 250

urban – non-urban comparisons from 12 species and 30 studies (i.e., the subset of

observations after 1992 for which geolocations were available).

Publication bias

We assessed the evidence for the existence of two types of publication biases, small-study and time-lag effects, following the recommendation in Nakagawa *et al.* (2022). We ran four additional multilevel meta-analytic models, two for lnRR and two lnCVR. Each of these models for lnRR and lnCVR included as a new moderator either the square-root of the inverse of the effective sample size or the mean-centered year of study publication (Trikalinos & Ioannidis 2005; Nakagawa *et al.* 2022). The amount of variation explained by these moderators (i.e., $R^2_{marginal}$) was calculated using the R function 'r2_*ml'* ('orchaRd' R package v.0.0.0.9000; Nakagawa *et al.* 2020). The statistical significance of these two moderators provides evidence for the existence of small-study and time-lag effects, respectively (Nakagawa *et al.* 2022).

Results

i) Is urbanisation associated with shifts in mean life-history avian traits?

Assessing the overall effect of urbanisation on phenotypic means across the three studied life-history traits, we found that urban populations tended to have overall lower mean values than their non-urban counterparts: life-history traits in urban habitats had, on average, 4.30% lower values than in non-urban habitats (lnRR mean estimate [95% Confidence Interval] = -0.044 [-0.103,0.015]; I^2_{total} = 98.6%; Figure S4; Table S1). 28.8% of the heterogeneity in lnRR was explained by phylogenetic and species-specific effects ($I^2_{phylogeny}$ = 4.3%; $I^2_{species ID}$ = 24.5%; Table S1), while 4.8% of heterogeneity was explained by consistent differences among studies (Table S1). Further analysis calculating urban effects per trait and accounting for potential covariation across the three investigated traits in the response to urbanisation (i.e., using a model with an unstructured variance-covariance matrix; see Methods and Table S2) confirmed that urban populations had lower mean values in every life-history trait: urban populations laid their eggs earlier (lnRR [95% CI] = -0.056 [-0.084, -0.028]; Figure 2a), laid

smaller clutches (lnRR [95% CI] = -0.075 [-0.109, -0.040]; Figure 2a), and tended to produce

fewer fledglings per clutch than non-urban populations (lnRR [95% CI] = -0.075 [-0.183,

0.034]; Figure 2a; Figure S5a). This meta-analytic model estimated different random effect

intercepts per trait and allowed for correlations across traits (see Methods for details). This

model revealed strong correlations in the response to urbanisation across traits: studies

reporting much earlier laying date in urban populations also reported more similar clutch size

and number of fledglings between habitats (Figure 3a & 3b). Likewise, studies reporting

large differences in clutch size between urban and non-urban populations also reported

large differences between habitats in number of fledglings (Figure 3c; see 'Study ID

(correlations)' in Table S3; i.e., correlations among studies in the values of lnRR for each

trait).

ii) Is urbanisation associated with changes in variation in avian life-history traits?

The coefficient of phenotypic variation in urban populations was, on average, 8.0% higher

than in non-urban populations (lnCVR mean estimate [95% CI] = 0.077 [-0.032, 0.186]; I^2_{total}

= 71.9%; Figure S6 and Table S1). Only 6.0% of the heterogeneity in lnCVR was explained

by phylogenetic and species-specific effects ($I^2_{phylogeny}$ = 3.0%; $I^2_{species\ ID}$ = 3.0%; Table S1),

while consistent differences between studies explained no heterogeneity in lnCVR ($I^2_{study\ ID}$ =

0.0%; Table S1). A subsequent model of lnCVR separating urban effects on phenotypic

variation per trait and accounting for potential covariation across the three investigated traits

in the response to urbanisation (see Methods and Table S4) revealed that the overall effect

of urbanisation on life-history trait variation was driven by urban populations having a more

variable phenology than their non-urban counterparts (lnCVR mean for laying date [95% CI]

= 0.193 [0.107, 0.279], i.e., 21.9% more variation, on average, in laying date in urban than

non-urban populations). Although the 95% confidence intervals overlapped zero, the

direction of the mean effects for clutch size and number of fledglings also reflected higher

phenotypic variation in urban populations compared to non-urban populations (Figure 2b;

lnCVR mean estimates [95% CI]: clutch size = 0.052 [-0.046, 0.149], number of fledglings =

0.064 [-0.071, 0.198]). We did not find evidence for the existence of correlations among life-

history traits in their lnCVR (Figure 3; models including correlations among traits scored

more than 1.57 AIC points below the top model, which only included independent random

intercepts per trait; Table S4). Differences in phenotypic variation between habitats were

quantitatively very similar when we meta-analysed the amount of total variation (i.e., lnVR;

Nakagawa *et al.* 2015) instead of the coefficient of variation (i.e., lnCVR; Figure S5).

iii) What is the temporal and spatial scale at which urbanisation affects phenotypic

variation?

Differences in phenotypic variation in laying date between the urban and non-urban habitat

arose from differences in variation within breeding seasons rather than between breeding

seasons (Table 1). While laying dates in urban populations were more variable than in non-

urban populations within breeding seasons (lnCVR mean estimate [95% CI] = 0.193 [0.094,

0.292]; Table 1), a subsequent meta-analytic model isolating effects on phenotypic variation

arising from between breeding season fluctuations revealed no strong difference between

urban and non-urban populations (lnCVR intercept mean [95% CI] = 0.007 [-0.133, 0.147];

Table 1). The sample size for this latter meta-analysis was almost four times smaller than for

the meta-analysis of within breeding season differences in variation; however, the estimated

lnCVR in both models showed large differences: the mean lnCVR within breeding seasons

was more than 27 times larger than mean lnCVR among breeding seasons (Table 1).

Furthermore, to assess whether urbanisation and/or habitat heterogeneity could explain

increased phenotypic variation in urban bird populations, we investigated the extent to which

our quantification of urban index and habitat heterogeneity predicted differences in

phenotypic variation across habitats. First, we confirmed that the urban populations included

in our meta-analysis were indeed exposed to higher levels of urbanisation than paired non-

urban populations (urban index in urban population ± SE = 0.641 ± 0.006; urban index in

non-urban population ± SE = 0.018 ± 0.006; at a spatial scale of 2000 m in both cases for

reference; Figure 4a). Including the difference in urban index and habitat heterogeneity

between paired urban and non-urban populations revealed that the more heterogeneous the

urban habitat was, the larger the phenotypic variation in this habitat compared to the non-

urban habitat; this effect was particularly strong at medium-large spatial scales (Figure 4c).

Differences in urban index between populations did not strongly explain variation in lnCVR

(Figure 4b).

iv) Assessment of publication bias

We found little evidence for the existence of small-study or time-lag biases in our data set.

Small-study effects and time-lag effects were not detected in our multilevel meta-analytic

models of lnRR (slope estimate for the square-root of the inverse of the effective sample size

[95% CI] = -0.024 [-0.187, 0.138], $R^2_{margnal}$ = 0.05%; slope estimate for year of publication

[95% CI] = -0.001 [-0.003, 0.002], $R^2_{margnal}$ = 0.05%). Small-study effects and time-lag effects

were not detected either in our multilevel meta-analytic models of lnCVR (slope estimate for

the square-root of the inverse of the effective sample size [95% CI] = 0.292 [-0.122, 0.705],

$R^2_{margnal}$ = 1.92%; slope estimate for year of publication [95% CI] = 0.001 [-0.004, 0.006],

$R^2_{margnal}$ = 0.11%). Overall, these findings suggest that our results do not suffer from

publications bias (Nakagawa *et al.* 2022).

504 Discussion

We compiled a global dataset of avian life-history traits for paired urban and non-urban

populations of the same species to assess how urban living is related to changes in mean

phenotypic values and variances of breeding phenology, reproductive effort, and

reproductive success. A phylogenetically multilevel meta-analysis of this dataset confirms a

well-documented effect of urbanisation on mean phenotypes: urban bird populations laid

earlier and smaller clutches than their non-urban counterparts. This model, however, also

reveals correlated responses to urbanisation across life-history traits: e.g., the earlier the

laying date in urban populations, the smaller the difference in clutch sizes between habitats.

Our study goes a step further than previous meta-analyses in urban ecology by explicitly

investigating how urbanisation could impact phenotypic variation. Our findings highlight that

urbanisation is not only associated with lower mean phenotypes, but also with increases in

phenotypic variation. Investigating the temporal and spatial scale at which increases in urban

phenotypic variation arise revealed hints at the ecological causes and evolutionary

consequences of increased phenotypic variation in urban bird populations.

Urbanisation has been associated with shifts in mean phenotypic values across a wide

range of organisms (Alberti *et al.* 2017; Merckx *et al.* 2018; Santangelo *et al.* 2022),

including birds, which show smaller body sizes and generally lower life-history trait values in

urban habitats (Chamberlain *et al.* 2009; Sepp *et al.* 2018; Thompson *et al.* 2022). Our

analyses expand the spatial, temporal and phylogenetic coverage of previous meta-analyses

of the avian literature (Chamberlain *et al.* 2009; Sepp *et al.* 2018), and agree on their

findings. Our results indicate that urban bird populations lay their eggs earlier and produce

smaller clutches, which results in a lower number of surviving nestling, than their non-urban

neighbouring populations. It should be noted, however, that our lnRR analysis indicates a

high heterogeneity. This finding indicates large variation among studies in how urbanisation

associates with changes in mean phenotypes and suggests that additional ecological traits

(e.g., diet or migratory strategy) may also affect how populations respond to urbanisation.

Our results also indicate that the mean response to urbanisation is correlated among traits.

Interestingly, we found that earlier laying dates in urban populations were associated with

smaller differences in clutch size and number of surviving nestlings between habitats. Clutch

size decreases with later laying date in several bird species (Rowe *et al.* 1994; Dunn &

Møller 2014), and these two traits (i.e., laying date and clutch size) are often hypothesized to

co-evolve because of a strong negative genetic correlation (Garant *et al.* 2008). All else

being equal, urban conditions triggering an earlier onset of reproduction (e.g., light pollution

(Dominoni *et al.* 2013) or increase resource availability during winter (Schoech *et al.* 2004))

could indirectly increase clutch size and, therefore, reduce differences in reproductive output

(clutch size and, possibly, number of fledglings) between urban and non-urban populations

that arise via other mechanisms (e.g., resource limitation in spring (Seress *et al.* 2018,

2020)). The extent to which mean phenotypic shifts represent adaptive responses to

urbanisation in birds, either via genetic changes or plasticity, or are maladaptive, is mostly

unknown (Lambert *et al.* 2020; see Santangelo *et al.* (2022) for evidence of adaptive

phenotypic shifts in urban plants). Our results, however, underscore that phenotypic shifts in

urban populations are widespread and that the response to urbanisation of associated life-

history traits should be investigated together.

Urbanisation has been recently hypothesised to increase phenotypic variation (Thompson *et*

al. 2022) and, indeed, higher variation in the morphological traits of urban great tits (*Parus*

major) and blue tits (*Cyanistes caeruleus*) has been recently reported (Thompson *et al.*

2022). Our findings greatly expand the evidence for this emerging hypothesis showing that

urbanisation is overall associated with increases in variation in laying dates across a wide

range of bird species. Previous studies have suggested that city characteristics, such as

warmer temperatures in early spring due to the urban heat island effect, could allow birds to

lay more clutches per season (Yeh & Price 2004; Schoech *et al.* 2008), with thereby longer

breeding seasons and hence higher phenotypic variation in urban laying dates (a similar

result has also been reported in Lepidoptera (Merckx *et al.* 2021)). This effect, however,

does not necessarily explain our results as our meta-analysis only included first clutch laying

dates per season. As such, our findings indicate that urban bird populations display more

variation in the onset of reproduction than their non-urban neighbours regardless of the total

number of clutches laid in urban and non-urban habitats.

Higher phenotypic variation in urban than in non-urban populations within breeding seasons

could be explained by at least two different evolutionary mechanisms, which could act in

isolation or in combination: differences in the underlaying additive genetic variance in laying

date, whereby urban birds have a wider range of genotypes for laying date; and / or differences in habitat heterogeneity influencing plasticity in laying date, whereby urban areas have larger environmental variation than non-urban habitat (Shochat *et al.* 2006; Heisler & Brazel 2018; Strubbe *et al.* 2020; Thompson *et al.* 2022). No study to date has investigated whether urban birds show higher additive genetic variance than non-urban populations. However, genetic analyses of European great tits in urban and non-urban habitats generally suggest small differences in the magnitude of genetic variation between habitats (Björklund *et al.* 2010; Caizergues *et al.* 2021; Salmón *et al.* 2021). This is, perhaps, not surprising given the high mobility of birds and the fact that gene flow between urban and non-urban bird populations likely occurs at a large spatial scale (Salmón *et al.* 2021). Interestingly, some studies have reported weaker selection for laying dates in urban areas than in non-urban habitats, suggesting relaxed selection on phenology in urban birds (Caizergues *et al.* 2018; Branston *et al.* 2021), which could increase the amount of phenotypic variation in phenology (Caizergues *et al.* 2018). Assessing differences in phenotypic variation between urban and non-urban populations of less mobile species will be important to evaluate how biological traits (e.g., dispersal ability) determine the evolutionary impact of urban ecological conditions. Previous work in mammal and amphibian species with lower dispersal ability than birds suggests a comparable level of (genetic) variation between urban and non-urban habitats (Fusco *et al.* 2021; Richardson *et al.* 2021).

Habitat complexity differs between urban and non-urban habitats (Arnfield 2003; Pickett *et al.* 2017). Our analyses indicate that differences in urban *versus* non-urban habitat heterogeneity could indeed contribute to explain the observed pattern of increased phenotypic variation in urban populations. Several ecological mechanisms could mediate this effect. Urban environments are characterised by an array of microhabitats with varying levels of human pressure, exotic plant species and resource availability. Thus, the intensity and timing of the environmental cues that birds use to time their reproduction could vary at a small local scale, increasing phenotypic variation in phenology in the presence of plasticity.

The existence of plastic responses to urban habitat heterogeneity, which our results might

indicate, do not preclude selection from acting on urban bird populations. First, plasticity is

an important mechanism of adaptation, sometimes aligned in direction with adaptative

genetic changes (De Lisle *et al.* 2022), and indeed often involved in adaptation to urban

environments (Halfwerk *et al.* 2019; Campbell-Staton *et al.* 2021). Second, plastic responses

can aid adaptation to urban conditions in the presence of genetic-by-environment

interactions by increasing the amount of genetic variation available for natural selection (Via

& Lande 1985). Addressing which evolutionary mechanisms cause the observed increase in

phenotypic variation in urban bird populations is beyond the scope of this study and we

acknowledge that these arguments are largely speculative at this point. However, our

findings highlight that eco-evolutionary processes may largely differ between urban and non-

urban bird populations and generate new avenues for future research in urban ecology and

evolution.

In agreement with our initial predictions, habitat heterogeneity was associated with the

magnitude of the difference in phenotypic variation between urban and non-urban bird

populations. However, we acknowledge that this analysis has a number of limitations and

that the results require cautious interpretation. First, only a subset of published studies

provided locations (i.e., coordinates) for their urban and non-urban study populations (30 out

of 65 published papers). When study site coordinates were provided, only one pair of

coordinates per study location were provided, making impossible to accurately assess the

actual area over which a given breeding population was studied. Additionally, it is common in

urban eco-evolutionary studies to monitor several populations within one single city.

However, in most studies, spatial information was provided at the scale of the whole city

(e.g., a single set of coordinates), preventing the accurate quantification of habitat

heterogeneity for every sub-population within a given urban habitat. These limitations

highlight that the ability to perform global meta-analyses on the effects of urban habitat

heterogeneity on phenotypic variation would be greatly improved if individual studies in

urban ecology provided accurate coordinates of the location of their study populations

(where ethical to do so). Reporting such information would allow future research synthesis to

quantify phenotypic variation within urban habitats (e.g., across different sub-populations in

the same city) as well as between urban and non-urban habitats.

Taken together, our results show that urbanisation is associated not only with changes of

mean phenotypic values, but also with increases in the amount of phenotypic variation that

avian populations display. Our analyses also highlight a temporal and spatial mechanism

that could generate such differences in phenotypic variation between urban and non-urban

habitats. We show that urban avian populations have a more variable phenology than non-

urban conspecifics within breeding seasons (i.e., differences in phenology across habitats

are seemingly not due to between-year fluctuations) suggesting that the ecological

conditions that generate such differences are constant across multiple years. Our coupled

spatial analysis indicates habitat heterogeneity and plastic responses as potential eco-

evolutionary mechanisms generating these results. The eco-evolutionary implications of

higher phenotypic variation in urban environments will likely vary among species (Thompson

et al. 2022) and our findings highlight the need for detailed investigation of these

consequences. To this end, long-term studies of individually marked individuals in replicated

paired urban and non-urban environments could be particularly fruitful to unravel whether

differences in phenotypic variation between urban and non-urban populations are caused by

differences in underlaying genetic variation and/or plastic responses to the urban

environment.

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Competing interests

The authors declare no competing interests.

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1050   **Tables**

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Table 1. Differences in variation (lnCVR) in life-history traits between urban and non-

urban populations at different temporal scales. Urban – non-urban differences in

variation (lnCVR) in laying date, clutch size and number of fledglings per clutch were meta-

analysed to assess differences in variation between urban and non-urban populations within

and among breeding seasons (e.g., different temporal scales). lnCVR estimates represent

meta-analytic model intercepts following the model structure presented in Table S5; positive

values indicate higher variation in urban populations than in non-urban populations and *vice*

versa. 'CI' = confidence interval; 'k' = sample size. Terms in italic bold highlight lnCVR

estimates whose 95%CIs do not overlap zero.

Temporal scale	lnCVR estimate [95% CI]			k
	Laying date	Clutch size	Number of fledglings	
Overall	*0.193*	0.052	0.064	409
	[0.107, 0.279]	[-0.046, 0.149]	[-0.071, 0.198]	
Within breeding season	*0.193*	0.024	0.149	376
	[0.094, 0.292]	[-0.110, 0.158]	[-0.025, 0.322]	
Among breeding seasons	0.007	0.077	0.011	101
	[-0.133, 0.147]	[-0.034, 0.189]	[-0.136, 0.158]	

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 Figures

Figure 1. Phylogenetic and geographical breadth of the meta-analytic data. (a) Phylogenetic tree of the 33 avian species included in the meta-analysis along with the number of effect sizes (i.e., urban – non-urban comparisons) included per species ('k'; which may encompass multiple years of study from the same publication) and the proportion of observations for each life-history trait (purple: Laying date; orange: Clutch size; Green: Number of fledglings). (**b**) Our meta-analysis included a broad range of species, as examples, left to right from top to bottom: *Sturnus vulgaris*, *Spinus tristis*, *Aphelocoma coerulescens*, *Strix aluco*, *Larus argentatus* and *Fulica atra*. All images are copyright free (CC - Public Domain Mark 1.0. Authors: Shenandoah National Park [first two images], Mike Carlo/U.S. Fish & Wildlife Service, Raini Svensson, Vanzan and Ekaterina Chernetsova) and were extracted from www.flickr.com. (**c**) Global map (excluding Antarctica) showing the location of each study included in the meta-analysis. Each point represents one study area

1076   in which one or more urban – non-urban pairs of populations were sampled across a varying

1077   number of years.

Figure 2. Urban populations have earlier phenology, lower reproductive output and more variable life-history traits than non-urban populations. (**a**) Urban populations laid earlier and had smaller clutches, producing fewer fledglings, than their paired non-urban populations (illustrated by negative lnRR estimates). (**b**) Our meta-analysis revealed that variation in life-history traits was higher in urban populations compared to non-urban counterparts, with a marked difference between habitats in laying date (illustrated by positive estimates of lnCVR). Model estimates for (**a**) lnRR and (**b**) lnCVR are shown along with their 95% confidence intervals per trait as calculated by our meta-analytic models accounting for correlated responses to urbanisation among traits (see Table S3 & Table S5 for full model outputs and Figure S4 and S6 for overall meta-analyses of lnRR and lnCVR). Number of studies included for laying date, clutch size and number of fledglings were 32, 42 and 44 studies respectively. Raw data and model estimates are presented in Figure S5. 'k' provides the number of urban – non-urban comparisons.

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Figure 3. Avian life-history traits show a correlated response to urbanisation.

Our meta-analysis allowed us to investigate correlated responses to urbanisation across the

three studied life-history traits (laying date, clutch size, number of fledgelings), and revealed

strong correlations in log response ratio (lnRR) but not log coefficient of variation ratio

(lnCVR). (**a**) Earlier laying dates in urban populations compared to non-urban counterparts

(i.e., negative values in the x axis) were associated with no differences in clutch size across

habitats (i.e., y axis values close to zero), defining a negative correlation between lnRR for

laying dates and clutch size. (**b**) A similar pattern was found between lnRR for laying dates

and number of fledglings, while (**c**) lnRR for clutch size and number of fledglings were

positively correlated (Table S2). (**d** - **f**) We found no statistical evidence for models including

correlations across traits in how urbanisation affected phenotypic variation (Table S5): (**d**)

differences between habitats in phenotypic variation in laying dates were not associated with

differences between habitats in phenotypic variation in clutch size or (**e**) number of

fledglings; and (**f**) differences between habitats in variation in clutch size were not associated

with differences between habitats in variation in number of fledglings. Points represent mean

values per study ± SE. Regression lines (mean ± SE) in **a** - **c** were fitted using linear

regressions to illustrate the correlation revealed by our trivariate meta-analysis (Table S3).

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Figure 4. Effects of habitat heterogeneity on the difference in phenotypic variation between urban and non-urban bird populations (i.e., lnCVR). (**a**) After quantifying urban index and habitat heterogeneity, we verified that urban populations had higher urban index (i.e., the proportion of landcover at a given spatial scale categorised as 'urban' [see methods]), which was indeed the case. The y axis represents the difference in urban index between urban and non-urban populations, with positive values representing urban populations with higher urban index than their non-urban neighbours. (**b**) Differences in urban index between urban and non-urban populations did not predict the magnitude of the difference in phenotypic variation between populations (i.e., lnCVR). This figure shows the estimated effect of differences in urban index between populations on lnCVR. Significant positive values indicate that the higher the difference in urban index between urban and non-urban populations, the higher the lnCVR value (i.e., larger values of phenotypic variation in urban populations compared to non-urban counterparts). (**c**) Differences in habitat heterogeneity between urban and non-urban populations did positively predict the magnitude of the difference in phenotypic variation between populations (i.e., lnCVR), particularly at large spatial scales. This figure shows the estimated effect of differences in habitat heterogeneity on lnCVR at different spatial scales. Significant positive values indicate that the higher the difference in habitat heterogeneity between urban and non-urban

1131 populations, the higher the lnCVR value (i.e., larger values of phenotypic variation in urban

1132 populations compared to non-urban counterparts). Points represent mean model estimates ±

1133 SE in **a**, and mean model estimates ± 95% confidence intervals (95%CI) in **b** and **c**. 'Spatial

1134 scale' refers to the radius of a circular area centred at each study location and over which

1135 urban index and habitat heterogeneity was calculated.