

Urban Signatures of Phenotypic Change: Emerging Patterns, Mechanisms, and Novel Hypotheses

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1 Humans challenge the phenotypic, genetic and cultural makeup of
2 species by affecting the fitness landscapes on which they evolve. Recent
3 studies show that cities might play a major role in contemporary evolution
4 by accelerating phenotypic changes in wildlife, including animals, plants,
5 fungi, and other organisms. Many studies of eco-evolutionary change
6 have focused on anthropogenic drivers, but none of these studies have
7 specifically examined the role that urbanization plays in eco-evolution or
8 explicitly examined its mechanisms. This paper presents new evidence
9 on the mechanisms linking urban development patterns to rapid
10 evolutionary changes for species that play important functional roles in
11 communities and ecosystems. Through a meta-analysis of experimental
12 and observational studies reporting more than 1600 phenotypic changes
13 in species across multiple regions, we ask whether we can discriminate
14 an urban signature of phenotypic change beyond the established natural
15 baselines and other anthropogenic signals. We then assess the relative
16 impact of five types of urban disturbances including habitat modifications,
17 biotic interactions, habitat heterogeneity, novel disturbances, and social
18 interactions. Our study shows a clear urban signal; rates of phenotypic
19 change are greater in urbanizing systems compared to natural and non-
20 urban anthropogenic systems. By explicitly linking urban development to
21 traits that affect ecosystem function, we can map potential eco-
22 evolutionary implications of emerging patterns of urban agglomerations
23 and uncover new insights for maintaining key ecosystem functions upon
24 which the sustainability of human wellbeing depends.

25 eco-evolution | urbanization | ecosystem function | sustainability

26 Introduction

27 Emerging evidence of phenotypic change on contemporary time-scales
28 challenges the assumption that evolution only occurs over hundreds or
29 thousands of years. Anthropogenic changes in ecological conditions can
30 drive evolutionary change in species traits that can alter ecosystem
31 function (1–3). However, the reciprocal and simultaneous outcomes of
32 such interactions have only begun to emerge (4). Despite increasing
33 evidence that humans are major drivers of micro-evolution, the role of
34 human activities in such dynamics is still unclear. Might human-driven
35 evolution lead to ecosystem change with consequences for human
36 wellbeing within contemporary time-scales (5, 6)?

37 To address this question, human-driven phenotypic change must
38 be considered in the context of global rapid urbanization. In 1950, 30
39 percent of the world's population lived in urban settlements (7). By 2014,
40 that figure had risen to 54 percent, and by 2050 it is expected to reach 66
41 percent (7). By 2030, urban land cover is forecast to increase by 1.2
42 million km², nearly tripling the global urban land area of 2000 and
43 potentially driving significant biodiversity loss (8). Urbanization drives
44 systemic changes to socio-ecological systems by accelerating rates of
45 interactions among people, multiplying connections among distant
46 places, and expanding the spatial scales and ecological consequences
47 of human activities to global levels (9).

48 A critical question for sustainability is whether, on an increasingly
49 urbanized planet, the expansion and patterns of urban environments
50 accelerate the evolution of ecologically relevant traits with potential
51 impacts on urban populations via basic ecosystem services such as food

52 production, carbon sequestration, and human health (e.g. infectious
53 diseases). In cities, subtle and perhaps not so subtle eco-evolutionary
54 changes are occurring at an unprecedented pace. Humans challenge the
55 phenotypic, genetic, and cultural makeup of species on the planet by
56 changing the fitness landscapes on which they evolve. Examples of
57 contemporary evolution (sometimes called rapid evolution) associated
58 with urbanization have been documented for many species (1, 5, 6, 10).

59 This paper examines the mechanisms linking urban development
60 patterns to contemporary evolutionary changes. Through a meta-
61 analysis of experimental and observational studies that report >1600
62 phenotypic changes in many species of birds, fish, arthropods, mammals,
63 plants, and algae across multiple regions, we investigated the emergence
64 of distinct signatures of urban-driven evolutionary change. We
65 hypothesize that shifts in the physical and socio-economic structure and
66 function of large urban complexes can drive rapid evolution on many
67 species that play important roles in communities and ecosystems. Thus,
68 urbanization-driven phenotypic change may, in turn, impact critical
69 aspects of ecosystem function.

70 We ask two questions:

71 [1] Is there evidence of an urban signature of phenotypic change
72 beyond the established natural and anthropogenic signals, accelerating
73 rates of phenotypic change in species across multiple regions?

74 [2] What are the relative impacts of five different types of urban
75 disturbance: habitat modification, biotic interaction, heterogeneity, novel
76 disturbance, and social interaction?

77 Our hypothesis is grounded in the growing evidence that
78 urbanization is a major driver of contemporary evolution through several
79 mechanisms that operate simultaneously across multiple scales. Urban
80 development changes habitat structure (i.e., loss of forest cover and
81 connectivity) and processes (i.e., biogeochemical and nutrient cycling),
82 and it also modifies biotic interactions (i.e., predation) (6). Humans in
83 cities also mediate eco-evolutionary interactions by introducing novel
84 disturbances and altering habitat heterogeneity. Urban environments can
85 facilitate hybridization by bringing together species that were previously
86 reproductively isolated (11). They can also isolate populations through
87 habitat fragmentation (12). In addition to changes in the physical

Significance Statement

Eco-evolutionary feedbacks on contemporary timescales were hypothesized over half a century ago, but only recently has evidence begun to emerge. The role that human activity plays in such dynamics is still unclear. Through a meta-analysis of experimental and observational studies that report >1600 phenotypic changes in species across regions and ecosystem types, we examine the evidence that the rate of phenotypic change has an urban signature. Our findings indicate greater phenotypic change in urbanizing systems compared to natural and non-urban anthropogenic systems. By explicitly linking urban development to trait changes that might affect ecosystem function, we provide insights into the potential eco-evolutionary implications for maintaining ecosystem function and the sustainability of human wellbeing.

89 template, humans in cities modify the availability of resources and their
90 variability over time, buffering their effects on community structure (12).
91 Complex interactions resulting from changes in habitat and biotic
92 interactions coupled with emerging spatial and temporal patterns of
93 resource availability might produce new evolutionary dynamics and
94 feedbacks. Furthermore, what makes cities unique is the pace of change
95 associated with increasing *social interactions* that amplify the direct and
96 indirect impacts of human agency, both locally and in distant places
97 (telecoupling). Understanding how urban-driven contemporary evolution
98 affects ecosystem functions and services will provide new insights for
99 maintaining biodiversity and achieving global urban sustainability.

100 Results

102 We discriminated the emergence of distinct signatures of
103 urbanization by statistically modeling an organism's phenotypic change
104 as a function of urban disturbances, urban proximity, and other previously
105 identified variables potentially affecting phenotypic change. Using
106 generalized linear mixed-effect models (GLMM), in an information-
107 theoretic framework to enforce parsimony and acknowledge model
108 uncertainty, we analyzed a modified and geo-referenced version of a
109 database of rates of phenotypic change that has been developed over
110 two decades (5, 13–17). After a series of quality filters, we retained for
111 analyses 89 suitable studies targeting 155 species, 175 study systems,
112 and >1600 rates of phenotypic change (Figure 1) (*Supporting Information*
113 *SI, p. 1-3*).

114 Statistical models including urban variables out-performed models
115 lacking urban variables, while accounting for anthropogenic context and
116 other putatively important variables described below. Hendry et al. (2008)
117 (15) showed that organisms in an anthropogenic context (e.g., pollution,
118 overharvest) had higher rates of phenotypic change compared to those
119 in a natural context. It was unclear whether urban variables would add
120 explanatory power after statistically controlling for the effect of

121 anthropogenic context. Our results showed that urban variables provide
122 substantial additional information for explaining phenotypic change, thus
123 warranting further consideration (*SI, p. 5-9*).

The multifarious effects of urban agglomerations occur across
multiple spatial scales (9). Hence, we evaluated the effects of
urbanization considering both variables determined by location relative to
urban agglomerations, and urban-driven processes regardless of
location. Urban predictor variables included *Urban Disturbance*
(categorical, 8 classes of urban-related mechanisms plus one reference
natural state), *City Lights* (ordinal, ranging from 0 to 1 – wildland to city),
Anthropogenic Biome (ordinal, ranging from 1 to 6 – dense settlements to
wildlands), and *Urbanization* (continuous, difference in *Anthropogenic*
Biome between years 1900 and 2000) (*SI, p. 4*). Since some phenotypic
changes were measured from populations at two locations (see *Design*),
continuous and ordinal predictor variables were calculated both as mean
and delta (difference) between the two samples underlying a phenotypic
change. Overall, this analysis included six urban variables representing a
range of possible mechanisms, scales, and proxies of urban drivers of
phenotypic change (see Materials and Methods). We also included three
unrelated variables that may affect rates of phenotypic change (13–15):
(1) number of *Generations* (continuous, log-transformed), (2) whether the
phenotypic change was estimated from a longitudinal or cross-sectional
study *Design* (categorical, 2 classes – allochronic and synchronic), and
(3) whether the phenotypic change had a demonstrated genetic basis or
not (labeled *GenPhen*, categorical, 2 classes – genotypic and
phenotypic). We refer to these three *a priori* variables (*Generations*,
Design, and *GenPhen*) as background variables.

We conducted exploratory multimodel ranking and inference based
on second-order Akaike Information Criteria (AICc) to evaluate the
relative ability of urban and background variables to statistically explain
the absolute magnitude of standardized phenotypic change, and to
assess effect sizes averaged over all possible models (18). A large model

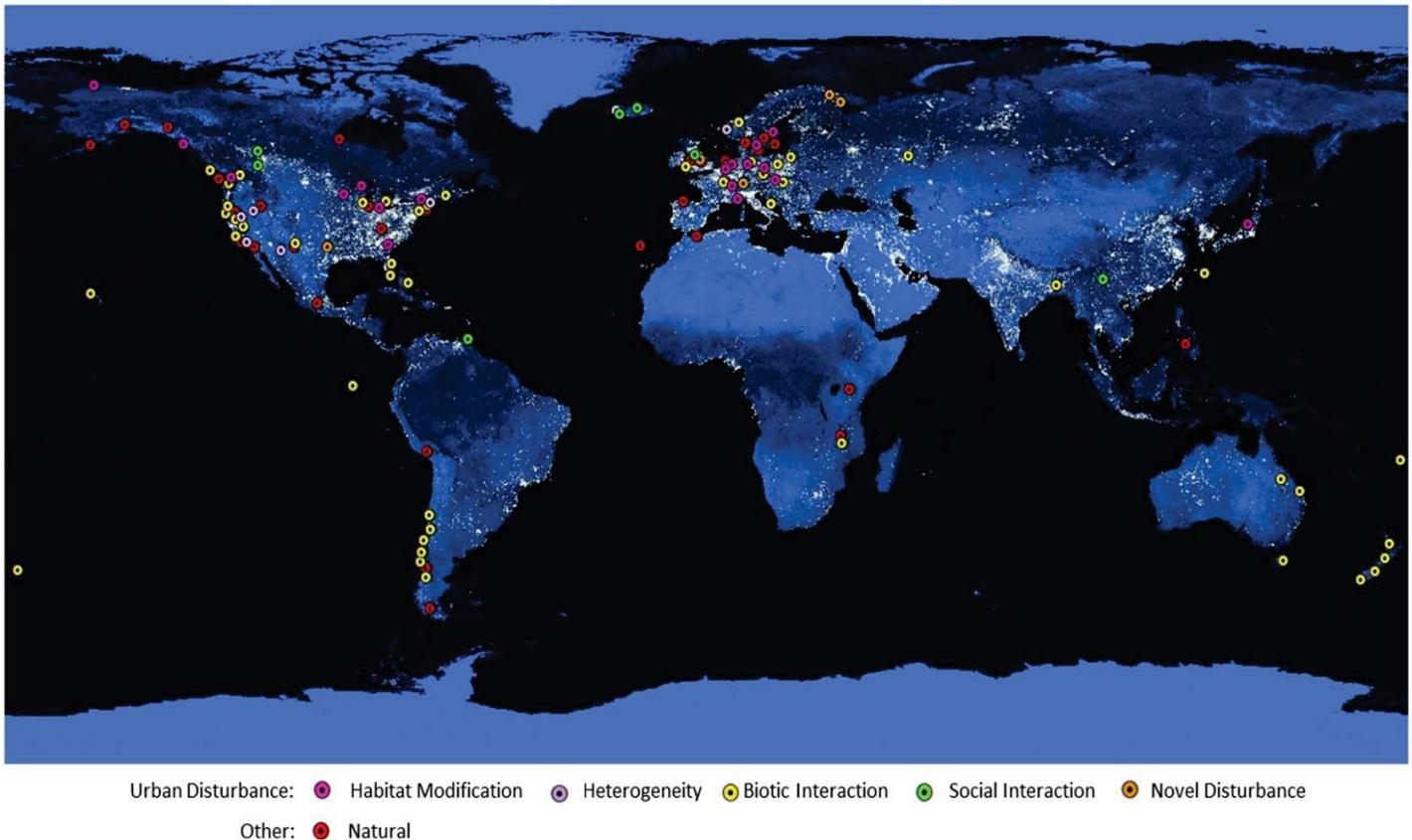


Fig. 1. Global distribution of study systems of trait changes in wild populations. Symbols represent Urban Disturbances, wherein each study system is categorized according to its primary driver of phenotypic trait change. White regions represent City Lights as interpreted by Zhou et al. (2015) (53). Background of the Earth in 2012 from NASA: earthobservatory.nasa.gov/Features/NightLights/page3.php

153 set (512 models) was created by considering all combinations of the nine
 154 explanatory variables in the fixed part of the GLMM (see Materials and
 155 Methods and *SI p.10-15*). The random part of all models was held
 156 constant, and included a random intercept per *Study System* to account
 157 for nested data structure, and a previously selected variance function that
 158 allowed the residual variance to scale with the expected response.
 159 Phenotypic change (square-rooted) was the response variable,
 160 measured as the absolute magnitude of phenotypic change standardized
 161 by character variation, a quantity known as Haldane numerator (19).

162 Top-ranked models consistently included urban variables. For
 163 example, the focal variables *Urban Disturbance*, *City Lights*, and
 164 *Anthropogenic Biome* are prevalent in top-ranked models and in the 95%
 165 confidence set, whereas, among background variables, *Generations* and
 166 *GenPhen* were prevalent (Figure 2). Model-averaged partial regression
 167 coefficients (incorporating model uncertainty) revealed several strong and
 168 many weak effects (Table 1). Among urban-related variables, phenotypic
 169 changes estimated from contrasts between urban vs. wild habitats (i.e.,
 170 Δ *City Lights*) were higher than contrasts within either urban or wild
 171 habitats. Mean *City Lights*, however, did not show a significant effect.
 172 *Urbanization*, inferred from land-cover change during the last century
 173 (mean *Urbanization*, with negative scores representing urbanization)
 174 showed a trend with highest rates of phenotypic change in urbanizing
 175 locations. This trend was not supported by the effect of contemporary land
 176 cover (*Anthrobiome*). *Urban Disturbance* had many effects. For example,
 177 social interactions, novel disturbances and introduction of predators,
 178 prey, hosts, or competitors, were associated with relatively high
 179 phenotypic change. Some effects were counterintuitive, e.g., habitat
 180 modification was associated with relatively low phenotypic change. The
 181 effects of *Urban Disturbance* were further illustrated by multimodel
 182 predictions made while the effects of other variables were statistically held
 183 constant (Figure 3). The range of effects attributed to Urban Disturbance
 184 on multi-model predictions were substantial compared to those of
 185 different combinations of background variables (*SI9*).

187 Discussion

188 Our results show a clear urban signal of phenotypic change and reveal
 189 variable effects of urban disturbance mechanisms. *Urban Disturbance*

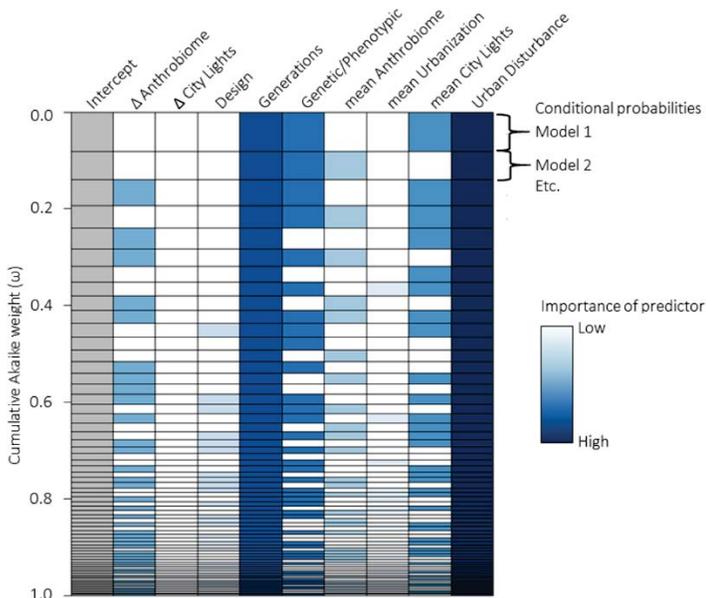


Figure 2. Representation of the AICc model-selection table. Rows represent models sorted by decreasing empirical support (row height represents model probability conditional on the full model set). Predictor variables were shaded if included in a model. Saturation corresponded to estimated variable relative importance. Note all high-ranked models contained urban (e.g., *Urban Disturbance*) and background variables (e.g., *Generations*). Models with little relative support were omitted for clarity (95% confidence set displayed).

190 drives changes in species traits that may play significant roles in the
 191 function of ecosystems and their stability, with significant consequences
 192 for sustainability. The large urban effect might be due to the multiple
 193 challenges that urbanization poses on adaptation. Multiple influences can
 194 increase the total strength of selection on a trait, or the number of traits
 195 under selection (20).

196 **Table 1:** Model-averaged coefficients from the full model set revealed
 197 several strong and many weak effects.

Parameter ^a	Estimate	Std. Error	Z-score	P-value
(Intercept)	0.545	0.097	5.619	0.000***
Generations	0.032	0.015	2.22	0.026*
Design – Synchronic	0.034	0.065	0.528	0.597
GenPhen – Phenotypic	0.104	0.052	2.006	0.045*
Mean. City Lights	0.056	0.033	1.695	0.090
Δ City Lights	0.072	0.031	2.344	0.019*
Mean Anthrobiome	-0.009	0.008	1.141	0.254
Δ Anthrobiome	-0.002	0.009	0.216	0.829
Mean Urbanization	-0.014	0.008	1.799	0.072.
U. Dist. – Hetero	0.019	0.097	0.200	0.841
U. Dist. – HabMod	-0.371	0.046	8.088	0.000***
U. Dist. – Novel	0.14	0.136	1.028	0.304
U. Dist. – Social	0.425	0.154	2.753	0.006**
U. Dist. – Int	0.077	0.073	1.061	0.289
U. Dist. – IntEco	0.345	0.126	2.738	0.006**
U. Dist. – Expalnt	0.005	0.09	0.059	0.953

198 Significance levels: P<0.001(***), P<0.01 (**), P < 0.05 (*), P < 0.1 (.)

199 ^a Abbreviations: Urban disturbance (U. Dist.), habitat modification
 200 (HabMod), heterogeneity (Hetero), and subcategories of biotic
 201 interaction: introduction (Int), introduction of predator/prey/
 202 host/competition (IntEco), range expansion after introduction (Expalnt).

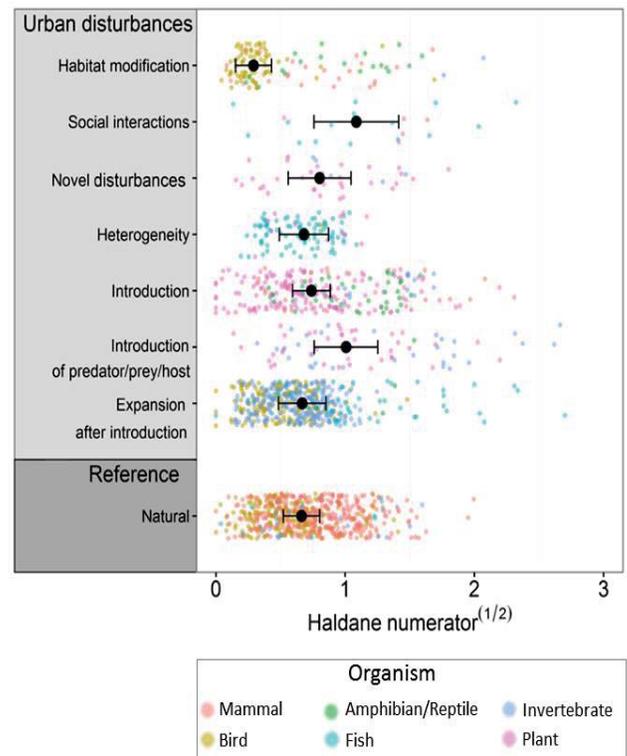


Figure 3. Multi-model predictions for *Urban Disturbance* categories. Some categories were associated with relatively high phenotypic change (i.e., *Social Interactions* and *Introduction of predator/prey/host/competitors*) whereas others associated with background or even stable phenotypes (*Habitat modification*). Modelled, but not shown variables were held constant at means (continuous variables) or reference values (categorical variables).

203	Urban Disturbance Mechanisms	269	generates unique system properties, and it can suppress the probability
204	<i>Urban Disturbance</i> represents coupled mechanisms through which urban	270	of ecological processes being controlled by a single taxon. Urbanization
205	development affects natural processes and evolutionary dynamics. Model	271	not only modifies existing disturbance regimes (e.g., fire and flood
206	predictions highlight three categories driving the urban signature: social	272	management); it also creates novel disturbances including disrupted
207	interactions, novel disturbances, and biotic interactions, specifically	273	dispersal pathways or stressors, eco-toxins and pollutants (33, 34).
208	introduction of predators, prey, hosts, and competitors. Anthropogenic	274	Examples of phenotypic changes in response to new toxins include
209	habitat modification was of lesser impact.	275	earthworms' tolerance to metals (33) and changes in endocrine systems
		276	of fish and birds (34).
210	<i>Habitat modification</i>		
211	Land cover conversion and loss of native habitat are major drivers	277	<i>Social Interactions</i>
212	of contemporary evolution. Surprisingly, habitat modification driven by	278	Perhaps the most significant quality that distinguishes cities is their
213	anthropogenic change was not among the dominant drivers of higher	279	pace of change. Urbanization changes the dynamics of socio-ecological
214	phenotypic change relative to the Natural context, suggesting that	280	interactions by increasing interactions among people (35), between
215	globally, natural drivers of contemporary evolution might have	281	people and other species (36), and among distant places (9). Important
216	greater effects on phenotypic traits. In part, the effect of urban land cover	282	properties of cities of all sizes (i.e., gross domestic product) increase, on
217	on phenotypic change is captured by <i>City Lights</i> . Yet urban-driven land	283	average, faster than city population size (superlinearity). Increasing social
218	conversion affects species composition through subtle mechanisms not	284	interactions accelerate environmental changes and human impact on
219	captured by <i>City Lights</i> , but captured by our <i>Urban Disturbance</i> variable.	285	phenotypic traits via selective harvest (5).
220	For example, changes in climate, artificial lighting, and availability of food	286	Socio-economic shifts associated with urbanization can explain the
221	are principal drivers of change in the timing and duration of reproduction	287	emergence of a detectable urban signature of evolutionary change. This
222	in some bird species (21). Changes in productivity—the rate at which	288	is also evident from the simultaneous effects of the <i>Urban Disturbance</i>
223	energy flows through an ecosystem— might explain species diversity	289	and <i>City Light variables</i> . The urban extent, regardless of boundary
224	patterns along the urban-to-rural gradient, though studies have produced	290	definitions, only partially defines the extent of urban disturbances. City
225	contradictory results. Changes in net primary production associated with	291	functions depend on highly interconnected infrastructures and on flows of
226	urban land-cover change affect faunal and plant species richness (22).	292	material, energy, and information from both proximate regions (e.g., via
		293	hydroelectric dams) and distant ones (e.g., via trade and
227	<i>Biotic Interactions</i>	294	telecommunication). In urbanizing regions, distant human-natural
228	We determined that among subcategories of biotic interactions,	295	interactions are currently more prevalent and faster (9), challenging the
229	introduction of predator, prey, host, or competition contributes to a higher	296	ability to disentangle urban vs. non-urban anthropogenic influences on
230	rate of phenotypic change compared to range expansion after	297	phenotypic traits.
231	introduction or introduction alone. Urban development creates new		
232	opportunities and challenges for species competition and predation, both	298	<i>Linking urban evolutionary change to ecosystem function</i>
233	as exotic species are introduced and as invasive species migrate in,	299	By explicitly linking urban development to traits that affect
234	taking advantage of poorly integrated communities and patches in the	300	ecosystem functions and services, we start to map the eco-evolutionary
235	urban setting. This might result in a colonization process, as more	301	implications of urban-driven phenotypic trait changes and identify existing
236	frequent introductions of exotic species translate into invasions (23). For	302	gaps in knowledge. The evolution of antibiotic resistance, pesticide
237	example, along a 140km urban-to-rural environmental gradient	303	resistance, host-pathogen co-evolution, and evolution in response to
238	originating in New York City, McDonnell et al. (1997) (24) found higher	304	harvest and habitat change provide examples of the link between human-
239	levels of earthworm biomass and abundance in urban forests compared	305	driven evolution and sustainability (51). Urban environments provide the
240	to rural ones, likely because of introduced species. Urbanization also	306	context for many such interactions, bringing people in to contact with
241	alters the way species distribute and interact (25). Marzluff (2005) (25)	307	novel pathogens, accelerating the spread of genes that confer resistance,
242	found that while diversity still emerges as the balance between extinction	308	and generally selecting for traits that enhance the survival and
243	and colonization, species invasion plays a prominent role.	309	reproduction of wild organisms within human-built environments. When
		310	this selection facilitates the survival of desirable species, it can have
244	<i>Heterogeneity</i>	311	important benefits for biodiversity, human health, and ecosystem
245	The results provide evidence that cities directly and indirectly affect	312	services. For example, evolution of <i>Daphnia</i> may help improve water
246	species diversity by altering spatial and temporal habitat heterogeneity.	313	quality in the face of cultural eutrophication (37). However, adaptation can
247	Increasing evidence supports the hypothesis that urban regions amplify	314	also enable the survival and reproduction of species that reduce
248	heterogeneity by the intensity and speed of human-biophysical and social	315	biodiversity and human wellbeing. For example, evolution of pesticide
249	interactions (26). Cities worldwide retain significant numbers of native	316	resistance in agriculture and antibiotic resistance in human medicine
250	species, but loss of functional heterogeneity driven by urbanization	317	represent critical threats to food security and human (38, 39). Thus,
251	affects niche differentiation and species diversity (27). Furthermore	318	evolution is instrumental to sustainability and should take on a prominent
252	changes associated with urban land uses act as filters in urban species	319	role in the future of sustainability science.
253	composition, and losses of native species drive the homogenization of	320	Table 2 identifies examples of traits for which there is evidence of
254	ecological structures and functions (28). Habitat patches and their	321	evolutionary response to environmental changes driven by urbanization.
255	ecological communities are often isolated by a matrix of built	322	The evolution of traits that control ecosystem processes could lead to
256	environments. Fragmentation of natural patches due to urbanization	323	significant changes in ecosystem functions (40). For example, primary
257	affects the diversity, structure, and distribution of vegetation, and	324	productivity is associated with consumers' traits that regulate their
258	movement of organisms (29). Change in temporal microclimatic	325	demands for resources. Evolution in such traits can affect nutrient cycling
259	heterogeneity (e.g., heat islands) can extend the growing season in	326	and ultimately the magnitude and spatial distribution of primary
260	temperate cities, and droughts in desert urban areas (30).	327	production (4). Seed dispersers have a significant impact on plant
		328	diversity and their functional roles in urban ecosystems. A great diversity
261	<i>Novel Disturbance</i>	329	of organisms modify the physical structure of estuarine and coastal
262	Novel disturbance emerges as a dominant driver of phenotypic	330	environments, particularly dune and marsh plants, mangroves,
263	change. Human-induced disturbances in urban environments maintain	331	seagrasses, kelps, and infauna (41). Evolution in ecosystem-engineering
264	urban habitats at an early successional stage (31). Such novel	332	traits has potential functional impacts on maintaining the stability and
265	disturbances alter resource availability, ecosystem productivity, and	333	resilience (e.g., flood control) of coastal cities and the capacity of cities to
266	species diversity (31). Cardinale et al. (2006) (32) suggested two ways	334	adapt to climate change.
267	that disturbance can moderate relationships between biodiversity and		
268	ecosystem functioning: it can increase the chance that diversity		

335 Table 2: Mapping urban-driven phenotypic trait change to ecosystem function. Documented phenotypic trait changes, urban drivers, and hypothesized
 336 eco-evolutionary feedback mechanisms.

Urban Signatures		Phenotypic trait	Eco-evolutionary feedback		Ref.
<i>Urban Disturbance</i>	<i>Mechanism</i>	Physiological	Ecosystem function	Feedback mechanism	
Novel	Exposure to effluent and associated extreme heat from a steam-electric plant	Mean heat coma temperature (thermal tolerance) in freshwater snails (<i>Physa virgata</i>)	Biodiversity	New 'physiological races'	(42)
Habitat modification	Anthropogenic acidification (pollution) of lake habitats	Survival in acidic conditions in moor frogs (<i>Rana arvalis</i>)	Biodiversity	Colonization	(21)
Novel	Electricity pylons create novel habitats characterized by high levels of zinc	Zinc tolerance in plants: <i>Agrostis capillaris</i> , <i>Agrostis stolonifera</i> , etc.	Nutrient cycling	Life history changes	(43)
		Morphological	Primary productivity	Consumer-resource dynamics	
Heterogeneity	Hydrological connectivity altered via a fish ladder	Body size in brown trout (<i>Salmo trutta</i>)	Biodiversity	Life history changes	(44)
Biotic interaction	Invasion of a molluskivorous crab (<i>Carcinus maenas</i>)	Shell thickness (mm) in periwinkle snail (<i>Littorina obtusta</i>)	Nutrient cycling	Trophic interactions	(45)
Social Interaction	Long-term selective harvesting of a medicinal plant	Size of American ginseng plants (<i>Panax quinquefolius</i>)	Primary productivity	Consumer-resource dynamics	(46)
		Behavioral	Biodiversity		
Biotic interaction	Introduction (translocation) of animals to a predator-free island	Antipredator behavior in multiple species of marsupials	Nutrient cycling	Allocation of time to foraging vs. vigilance	(47)
		Phenological/Life History	Primary production	Consumer-resource dynamics	(48)
Heterogeneity	Temporal heterogeneity in water availability (drought)	Flowering time in Field mustard (<i>Brassica rapa</i>)	Biodiversity	Colonization	(49)
Habitat modification	Global warming	Seasonal onset of reproduction in 65 species of migratory birds	Biotic control	Novel competition	

337

338 **Conclusions**

339 Rapid urbanization poses new challenges for species—some will
 340 go extinct (at least locally) while others will adapt or relocate (50). Our
 341 paper asked whether we can detect signatures of eco-evolutionary
 342 change across taxa in urban ecosystems and determine the extent to
 343 which systemic changes to socio-ecological systems associated with
 344 urban agglomerations might accelerate phenotypic change. We found a
 345 clear urban signal of phenotypic change, and greater phenotypic change
 346 in urbanizing systems compared to natural and non-urban anthropogenic
 347 systems. By explicitly linking urban development to heritable traits that
 348 affect ecosystem function, we can begin to map the eco-evolutionary
 349 implications of human-induced trait changes for Earth's evolution.

350 We posit that urban-driven contemporary evolution will affect
 351 sustainability, from the level of the urban ecosystem to the planetary
 352 scale. We suggest that conservation biologists should pay increased
 353 attention to mechanisms by which the emergent human habitat influences
 354 population persistence (4). Such understanding will provide new insights
 355 for maintaining ecosystem function in the long term and can direct policy
 356 makers toward novel sustainability solutions (51).

357 **Materials and Methods**

358 Database on rates of phenotypic change
 359 We improved an existing database on rates of phenotypic change (5, 13–17). We added
 360 new data published up to August of 2015. Studies were surveyed by searching ISI Web of
 361 Science, Google Scholar, and cross-references, using *ad hoc* keywords (e.g., quantitative
 362 trait, evolutionary change, rapid evolution, eco-evolutionary, anthropogenic change, urban
 363 disturbances, and system stability). Studies were screened for eligibility (see *Database*
 364 *filtering in SI*), and, if selected, phenotypic rates were extracted (see *Statistical analyses*)
 365 and classified according to qualities of the study system including ecological and
 366 anthropogenic contexts (5). Each row corresponded to one phenotypic change rate estimate
 367 and associated contextual attributes including type of study: *allochronic* for longitudinal
 368 studies, or *synchronic* for cross-sectional studies comparing samples obtained
 369 synchronously from populations derived from a common ancestral population. Rates were
 370 classified according to if phenotypic change could be attributed to quantitative genetic
 371 effects (*Genetic*), or could not be distinguished from phenotypic plasticity (*Phenotypic*).

372 Generations was calculated as the number of years between population samples (or since
 373 population divergence, if *Synchronic*) divided by expected generation time.

374 The data set had a hierarchical structure, with variable numbers of phenotypic
 375 change estimates (from different morphological characters and or populations) within study
 376 systems, species, and taxa. Study system was defined as (a) population(s) of a species
 377 within a geographical region putatively exposed to similar environmental effects and high
 378 gene flow potential. Two important contextual variables were evaluated at the study-system
 379 level: whether study systems were regarded as evolving in an anthropogenic or a natural
 380 *Context* (15), and *Urban disturbance* (see next section).

381 **Urban Disturbance classification and geo-referencing**

382 We use the global urban area map at 1 km spatial resolution developed by Zhou et al. (2015)
 383 (52). The map is based on a cluster-based method to estimate optimal thresholds for
 384 mapping urban extent using DMSP/OLS NTL to account for regional variations in urban
 385 clusters, calibrated using high-resolution land cover data for the US and China in 2005 (53).
 386 The anthropogenic biome of all samples was based on the Anthropogenic Biomes
 387 geodataset for the year 2000 (54).

388 For samples in study systems in which the driver of evolutionary change is
 389 anthropogenic, we classify the *Urban Disturbance* as social interaction, biotic interaction
 390 (with subcategories for introduction, introduction of predator/prey-host/competition, and
 391 range expansion after introduction), habitat modification, heterogeneity, or novel
 392 disturbance (6) (See *Urban disturbance* classification in SI). Habitat modification represents
 393 changes in climate, modification of the landscape, or pollution. Biotic interactions stem from
 394 introductions, and are subcategorized depending on the study organism's ecological role:
 395 introduced species vs. species in its native range responding to an introduction. Introduced
 396 species are further divided into species adapting in a new range following introduction vs.
 397 introduced species adapting following range expansion. Heterogeneity can refer to
 398 heterogeneity in space or time, e.g., pulsed disturbances from discharge of a geothermal
 399 plant (55). Novel disturbances require novel adaptations, e.g., rapid evolution of zinc
 400 tolerance (43). Social interactions refer to direct or intentional results of human agency, e.g.,
 401 trophy hunting (56).

402 **Statistical analyses**

403 We used an information-theoretic approach to rank statistical models and conduct
 404 multimodel inference, based on AICc (57)(61, 23). AICc favors model fit (minimizing

deviance) while avoiding model overfitting (penalizing for the number of estimated parameters, K), and was the basis for enforcing the parsimony principle given our finite sample sizes (1663 rates nested in 175 Study systems). The statistical models were generalized linear mixed-effect models (GLMM). The response variable, phenotypic change (square-root transformed), was measured as the absolute magnitude of phenotypic change standardized by character variation (Haldane numerator; 19, as formulated in 13). The square-root transformation minimized patterns in adjusted residuals plots in preliminary analyses. Because the data had a hierarchical structure, study system was always modelled as a random effect, with combinations of background and urban variables (fixed effects):

$$H^{(1/2)}_i = \alpha_{j(i)} + \beta X_i + \mathcal{E}_i$$

$$\alpha_j \sim N(\mu, \sigma^2 \alpha)$$

$$\mathcal{E} \sim N(0, f(\gamma))$$

where the indexes i run from 1 to number of observations, and j run from 1 to number of study systems, $H^{(1/2)}$ is the response variable (square-root of Haldane numerator), α is normally distributed with mean μ (overall intercept) and variance $\sigma^2 \alpha$, allowing for varying intercepts per study system, β a vector of partial regression coefficients related to a matrix of explanatory variables X , and \mathcal{E} , the residual error with variance γ , which was modelled as follows:

$$\gamma = 0.1494 * (C + |\text{fitted}|)^P$$

where C is a constant by stratum (0.3233 for genetic; 0.1249 for phenotypic), and P an exponent of absolute fitted values by stratum (2.0754 for genetic; 1.2376 for phenotypic). Hence, the chosen residual variance increased exponentially with fitted values, and slightly more so in genetic than phenotypic rates (*SI*, p. 10-15).

We used exploratory multimodel inference to assess the relative importance of predictor variables (urban or not) for phenotypic change, and to make predictions about contrasting urban-related scenarios that considered information contained in all models. From 3 background plus 6 urban variables, we combined 9 predictor variables to form $2^9 = 512$ models, including a null model (intercept only), and excluding interactions. All models were fitted through maximum likelihood (ML) in the R package nlme (58). Models were ranked according to decreasing values of AICc (59), and further evaluated using standard methods after re-fitting through REML estimation (60). Predictor variable relative importance was calculated by the sum of the Akaike weights of all models containing a particular predictor variable. Similarly, model-averaged partial regression coefficients were Akaike-weighted averages of coefficients from all models containing a particular term (18). Model ranking and inference was conducted in the R package MuMin v1.15.6 (57) (*SI*, p.10-15).

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