



1 **Biotic factors dominantly determine soil inorganic carbon stock across**

2 **Tibetan alpine grasslands**

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14



15 **Abstract.** Soil inorganic carbon (SIC) pool is a major component of soil C pools, and
16 clarifying the predictors of SIC stock is urgent for decreasing soil C losses and
17 maintaining soil health and ecosystem functions. However, the drivers and their relative
18 effects on the SIC stock at different soil depths remain largely unexplored. Here, we
19 conducted a large-scale sampling to investigate the effects and relative contributions of
20 abiotic (climate and soil) and biotic (plant and microbe) drivers on the SIC stock
21 between topsoils (0–10 cm) and subsoils (20–30 cm) across Tibetan alpine grasslands.
22 Results showed that the SIC stock had no significant differences between the topsoil
23 and subsoil. The SIC stock was positively associated with altitude, pH, and sand
24 proportion, but negatively correlated with mean annual precipitation, plant
25 aboveground biomass, plant coverage, root biomass, soil available nitrogen, microbial
26 biomass carbon, and bacterial and fungal gene abundance. For both soil layers, biotic
27 factors had larger effects on the SIC stock than abiotic factors did. But the relative
28 importance of these determinants varied with soil depth, with the effects of plant and
29 microbial variables on SIC stock weakening with soil depth, whereas the importance of
30 climatic and edaphic variables increasing with soil depth. Specifically, bacterial and
31 fungal gene abundance and plant coverage played dominant roles in regulating SIC
32 stock in the topsoil, while soil pH contributed largely to the variation of SIC stock in
33 the subsoil. Our findings highlight differential drivers over SIC stock with soil depth,
34 which should be considered in biogeochemical models for better simulating and
35 predicting SIC dynamics and its feedbacks to environmental changes.

36



37 **1 Introduction**

38 Soils store approximately 1,500 Pg of organic carbon (SOC) and 940 Pg of inorganic
39 carbon (SIC) to a depth of 1 m (Batjes, 1996; Jobbágy & Jackson, 2000), which are the
40 largest carbon (C) pool in the terrestrial ecosystem and play a critical part in the global
41 C cycling (Darwish et al., 2018; Lal 2004; Prietzel et al, 2016). Compared to the
42 relatively short turnover time of SOC, SIC has a long residence time due to soil
43 weathering (Monger et al, 2015; Zang et al, 2018), which is considered to be fairly
44 stable and has less contribution to changes in terrestrial ecosystem C balance (Yang et
45 al, 2012). Therefore, previous studies have paid little attention to SIC. However, recent
46 studies suggest that SIC is also responsive to anthropogenic activities and global
47 climate changes such as soil acidification, atmospheric N deposition, and global
48 warming (Yang et al, 2010; Song et al, 2022), acting as a critical C source (Liu et al,
49 2020) or C sink (Gao et al, 2018; Liu et al, 2021). Thus, the preservation of SIC and its
50 roles in climate mitigation should not be neglected, especially in arid and semi-arid
51 grasslands where store a large amount of SIC (Yang et al, 2012).

52 SIC stock and stability can be fundamentally altered by an array of abiotic and biotic
53 processes (Raza et al, 2020). High precipitation can promote soil silicate minerals
54 weathering and removal of base cations (Ca^{2+} , Mg^{2+} , K^{+} , and Na^{+}) by leaching (Vicca
55 et al, 2022). Soil acidification due to atmospheric nitrogen (N) and acid deposition and
56 the nitrification of NH_4^{+} may greatly accelerate soil carbonate dissolution and CO_2
57 releases (Raza et al, 2020; Song et al, 2022). Plant growth can deplete soil carbonates
58 by releasing proton and organic acids from root rhizosphere (Goulding et al, 2016;



59 Kuzyakov & Razavi, 2019), and biological N₂ fixation by some legumes are likely to
60 cause SIC losses (Tang et al, 1999). Furthermore, plant autotrophic and microbial
61 heterotrophic respiration often facilitate carbonate dissolution by enhancing CO₂ partial
62 pressures (An et al, 2019; Liu et al, 2021). Nevertheless, how these abiotic and biotic
63 factors affect SIC stock and what is the relative importance of these confounding drivers
64 remain largely uncertain.

65 Previous studies on SIC stock mostly have focused on the topsoil, while the patterns
66 of SIC stock in the subsoil on a large scale remain elusive. The predictors of SIC stock
67 in the subsoil may differ from those in the topsoil due to distinct soil microenvironments,
68 soil physicochemical properties, root exudates, and microbial abundance and functions
69 (Jia et al, 2017). For instance, the topsoil has larger root biomass and higher microbial
70 activity than the subsoil, but the subsoil tends to preserve soil parent material because
71 of the weakened weathering by the isolation of heat and energy from the surface soil
72 (Crowther et al, 2016). Thus, the abiotic and biotic variables may exhibit different
73 effects on SIC stock in the subsoil compared to the topsoil due to the various importance
74 of these variables.

75 The Tibetan Plateau has the largest alpine grassland on the Eurasian continent,
76 which is a vital component of global terrestrial ecosystems, providing an ideal platform
77 to explore SIC stock and its determinants (Wang et al, 2002; Yang et al, 2010). During
78 the past several decades, the plateau has experienced significant warming (Wang et al,
79 2008) and pronounced atmospheric N deposition (Liu et al, 2013; Yu et al, 2019). This
80 continuous warming and N deposition have resulted in a significant increase in plant



81 growth and soil acidification (Ding et al, 2017; Yang et al, 2012), which could be likely
82 to induce potential CO₂ releases from soil carbonates by biogeochemical process (Raza
83 et al, 2020). However, a general understanding of SIC stock with soil depth across
84 Tibetan alpine grasslands remains unexplored. Here, we researched the relative
85 importance of climatic, edaphic, plant, and microbial variables to SIC stock at different
86 soil layers along an approximately 3,000 km transect of alpine grasslands on the Tibetan
87 Plateau, spanning a broad range of climatic and geographical conditions. Specifically,
88 two key questions are addressed in this study: (1) what are the differences of SIC stock
89 between the topsoil and subsoil? (2) how does the relative importance of climatic,
90 edaphic, plant, and microbial variables to the variation of SIC stock along with soil
91 depth?

92



93 2 Material and methods

94 2.1 Study area and field sampling

95 During July, August, and September 2020, we conducted large-scale systematic field
96 surveys and samplings in Tibetan alpine grasslands. The total 25 sampling sites covered
97 approximately 3,000 km and included three grassland types (i.e, 11 alpine meadow, 8
98 alpine steppe, and 6 alpine desert sites). The distance between nearby sampling sites
99 was about 120 km. The study sites cover a broad geographic and climatic range, with
100 longitude and latitude ranging from 79°49'39" to 102°25'31" E and 31°06'37" to
101 32°43'09" N, respectively, and the altitude ranging from 3500 m to 5016 m. These sites
102 covered a broad precipitation gradient varying between 72 mm and 706 mm. The mean
103 annual temperature (MAT) ranged from -3.9°C to 5.8°C. The plant communities were
104 dominated by *Kobresia tibetica Maxim*, *Stipa caucasica*, *Kobresia pygmaea*, *Stipa*
105 *purpurea*, and *Leontopodium pusillum*. Soils were *Cambisol* and some were loess-
106 derived *Luvisol*. The site location, grassland type, climatic, and plant parameters were
107 detailed in Table S1.

108 2.2 Climatic data

109 The climatic data were derived from the LPSDC (Loess Plateau Scientific Data Center,
110 <http://loess.geodata.cn/>) (Peng et al, 2019). The Kriging interpolation was conducted to
111 obtain spatial distributions of 30-year MAT and MAP (1987-2017) at each sampling
112 site by a geographic coordinate system.



113 **2.3 Soil properties**

114 At each site, we selected four 1 m × 1 m plots for soil and plant samplings and the
115 distance between nearby sampling plots was 25 m. In each plot, a 7.5-cm diameter soil
116 drill was used to take five soil cores at fixed soil depths (0-10 cm, 10-20 cm, and 20-30
117 cm), and a 2-mm mesh was used to remove stones. We used soil samples from 0–10 cm
118 and 20–30 cm to represent the topsoil and subsoil, respectively, according to previous
119 studies (Angst et al, 2021; Rumpel & Kögel-Knabner 2011; Zhou et al., 2021). After
120 mixing, 100 g of fresh soils from each plot were collected and stored in a –4°C portable
121 icebox, then returned to the laboratory and stored at –20°C for microbial properties.
122 The rest soil samples about 700 g were also sent back to the laboratory and air-dried for
123 measurements of other soil properties. A 40 cm × 40 cm × 40 cm (length × width × depth)
124 pit was dug for measuring soil bulk density (BD) by using a constant volume soil
125 sampling drill (100 cm³), and the undisturbed soil was preserved in aluminum specimen
126 boxes returning to the laboratory and oven-dried for 48 hours at 105°C and weighed.
127 The oven-dried soil (20 g) was screened into gravel by sifting through a 2-mm mesh
128 sieve and gravels larger than 2 mm were collected and weighed to determine the
129 percentage of gravels. Soil pH (1:25 soil: H₂O) was measured using a soil pH meter,
130 and available nitrogen (AN) was determined by the alkaline-hydrolysis diffusion
131 method. A laser particle analyzer (Mastersizer 2000, Malvern Panalytical, UK) was
132 applied to measure soil mechanical compositions, including clay (< 2 μm), silt (2-50
133 μm), and sand (> 50 μm) proportion. SIC was determined by using an inorganic C
134 analyzer (multi EA® 4000; Analytic Jena, Germany). The multi EA 4000 C elemental



135 analyzer was equipped with the automatic TIC solids module and calibrated before the
136 analysis. The sample boat was acidified automatically with 40 % H₃PO₄ in the reactor
137 of the TIC module. And the CO₂ from the carbonate was released, the measuring gas
138 was dried and cleaned and the carbon content was measured by means of the wide-
139 range NDIR detector. Before being analyzed directly, all soil samples were ground into
140 solid fine powders with a mortar, and for the determination of TIC, a standard, prepared
141 by solids-dilution of CaCO₃ with SiO₂ (0.2 % C), was used, with weighting range 7-200
142 mg, to cover a wide concentration range.

143 **2.4 Plant properties**

144 In each plot, we estimated plant coverage (PC) by the projection method, namely the
145 proportion of vegetation projection to the area of the sampling plot. In addition, plant
146 aboveground biomass and belowground roots were clipped and collected, respectively,
147 then oven-dried at 60°C and weighed to determine plant aboveground biomass (PAB)
148 and root biomass (RB).

149 **2.5 Microbial attributes**

150 Soil microbial biomass carbon (MBC) was measured by using a chloroform
151 fumigation-extraction procedure (Brookes et al, 1985). Briefly, 10 g of unfumigated
152 and chloroform-fumigated fresh soil samples were extracted by using 0.5 M K₂SO₄
153 after 24 h of incubation, respectively. Then, the extracts were analyzed by using a TOC
154 analyzer (multi N/C® 3100; Analytic Jena, Germany). The MBC was determined by



155 the differences in C concentrations between unfumigated and chloroform-fumigated
156 samples, and the correction factor (i.e, $KC = 0.45$) was used to convert microbial C to
157 MBC (Joergensen, 1996).

158 Real-time polymerase chain reaction (qPCR) was used to quantify bacterial (BA)
159 and fungal gene abundance (FA) by the absolute quantification method based on the
160 gene copy number (Tatti et al, 2016). Each reaction was carried out 3 times with a
161 mixture of a total 20 μL volume, including 2 μL of DNA template, 10 μL of 2 \times ChamQ
162 SYBR Color qPCR Master Mix, and 0.4 μL (5 μM concentration) each of forward and
163 reverse primer specific for each gene. And the PCR conditions were 95 $^{\circ}\text{C}$ for 5 min,
164 then 40 cycles for the 18S rRNA gene and 16S rRNA gene. Each cycle involved melting
165 at 95 $^{\circ}\text{C}$ for 30 s, annealing at 55 $^{\circ}\text{C}$ for 30 s, an extension of 72 $^{\circ}\text{C}$ for 40 s, and finally
166 10 $^{\circ}\text{C}$ until terminated. And the primer pair SSU0817/1196 and Eub338/Eub806 were
167 used for amplifying fungi and bacteria in PCR amplification, respectively. Then the
168 DNA concentration was determined by using a QuantiFluorTM-ST fluorescent
169 quantitative system (Promega, Fitchburg, WI, USA). The abbreviations of all variables
170 were detailed in Table S2.

171 2.6 Statistical analyses

172 The total SIC density (C stock per land area) in each soil depth layer was calculated
173 using Equation (1) (Pan et al, 2019):

$$174 \text{ SIC density (g C m}^{-2}\text{)} = \text{SIC (g C kg}^{-1}\text{)} \times \text{BD (g cm}^{-3}\text{)} \times \text{d (cm)} \times (1-g) / 100 \quad (1)$$



175 where SIC is soil inorganic C content, d is the depth of the soil layer (0.1 m), BD is
176 bulk density, and g is the percentage of gravel fraction (>2 mm).

177 First, the differences of SIC stock and corresponding abiotic and biotic variables
178 between the topsoil and subsoil were examined by T -test. Second, SIC density and
179 various abiotic and biotic variables were log-transformed and standardized (z-score
180 normalization) to perform the assumption of normality and homogeneity by Shapiro-
181 Wilk and Levene's test, respectively (Pan et al, 2021). Then the linear regressions were
182 used to test SIC density about different variables for both the topsoil and subsoil across
183 sites.

184 Third, a linear model was employed to examine SIC density with abiotic and biotic
185 variables by using the maximum likelihood estimation with the `lm` package. And the
186 relative effect of the parameter estimates was calculated to evaluate the relative
187 importance of drivers controlling SIC density. Also, SIC density and abiotic and biotic
188 variables were standardized before analyses, using the Z-score to interpret variable
189 estimates on a comparable scale (Gross et al, 2017).

$$190 \text{Log (SIC density)} = \beta_0 + \beta_1 \log X_1 + \beta_2 \log X_2 + \dots + \beta_{12} \log X_{12} \quad (2)$$

191 where β_0 and β_i ($i=1, 2, 3 \dots 12$) are intercept and coefficients, respectively. To
192 explore the determinants of SIC density in different soil depths across all sites, the
193 absolute values of slopes of the variables were extracted and plotted. Then, 12
194 controlling variables were categorized into four groups, including climatic (MAP, MAT,
195 and altitude), edaphic (pH, AN, and sand proportion), plant (PB, PC, and RB), and



196 microbial (MBC, BA, and FA) factors, to quantify their relative contribution to SIC
197 density (Fang et al, 2019).

198 Furthermore, the relative importance of abiotic (climatic and edaphic) and biotic
199 (plant and microbial) variables in determining SIC density was quantified by
200 performing variation partitioning analyses (VPA) by using the “vegan” package in R
201 4.1.3.

202



203 **3 Results**

204 **3.1 SIC density and influencing variables in different soil depths**

205 SIC density and SIC content had no significant differences between the topsoil and
206 subsoil, but bulk density in the subsoil was much higher compared with the topsoil.
207 Specifically, SIC density in the topsoil and subsoil ranged from 1.8 g C m⁻² to 3271 g
208 C m⁻² and 5.4 g C m⁻² to 3214 g C m⁻² across 25 sampling sites, with an average of 802
209 ± 220 g C m⁻² and 814 ± 236 g C m⁻², respectively (Fig. 1). No significant changes in
210 SIC density with soil depth were observed in both the alpine steppe and alpine desert
211 ($p=0.113$ and $p=0.068$, respectively; Fig. 1), but SIC density was higher in the subsoil
212 than that in the topsoil in the alpine meadow ($p = 0.002$, Fig. 1).

213 Meanwhile, the majority of abiotic and biotic drivers had significant differences
214 between the topsoil and subsoil (Table 1). RB, AN, MBC, BA, and FA in the topsoil
215 were significantly larger than those in the subsoil (all $p<0.001$). In contrast, pH was
216 significantly lower in the topsoil than in the subsoil ($p<0.001$, Table 1). However, the
217 sand proportion between the two soil depths had no significant differences (Table 1).

218 **3.2 Associations of SIC density with abiotic and biotic variables**

219 The SIC density was closely related to multiple abiotic and biotic variables (Figs 2 and
220 3). For both the topsoil and subsoil, the SIC density was positively associated with
221 altitude, pH, and sand proportion, but negatively correlated with MAP, PAB, PC, RB,
222 AN, BA, and FA. The SIC density showed a negative correlation with MBC in the



223 topsoil (Fig. 2), but not in the subsoil (Fig. 3). Meanwhile, the SIC density in both two
224 soil depths did not correlate with MAT (Figs. 2 and 3).

225 **3.3 Determinants of SIC density in different soil depths**

226 The linear model and VPA collectively displayed that the predominant drivers of SIC
227 density differed with soil depth (Figs. 4 and 5). Specifically, for the topsoil, the linear
228 model revealed that microbial and plant variables largely explained the variations in the
229 SIC density, followed by edaphic variables and climate contributed the least (Fig. 4).
230 Among these variables, PC, BA, and FA exhibited larger effects on the SIC density
231 compared with other controlling factors (Fig. 4). Also, the VPA analysis illustrated that
232 biotic factors explained the majority variation of SIC density compared with abiotic
233 factors (Fig. 5). For the subsoil, the linear model showed that edaphic variables largely
234 explained the variation in SIC density, followed by microbial and plant variables, and
235 climate contributed the least (Fig. 4). Among these variables, the soil pH had larger
236 contributions to the variation of SIC density rather than others (Fig. 4). Meanwhile, the
237 VPA analysis confirmed that the effects of biotic factors on SIC density were larger
238 than those of abiotic factors in the subsoil (Fig. 5).

239



240 **4 Discussion**

241 To the best of our knowledge, this study was the first to afford large-scale evidence of
242 the relative contribution of abiotic and biotic drivers to the variation of SIC stock at
243 different soil depths, which has considerable implications for grasping the importance
244 of SIC in the ecosystem C cycling. Since considerably stable characteristics and the
245 long turnover time (Mi et al, 2008; Yang et al, 2010; Zamanian et al, 2018), SIC stock
246 is traditionally considered to be dominated by abiotic factors including soil moisture,
247 soil pH, CO₂ partial pressure, and Ca²⁺ concentrations according to the equilibrium of
248 carbonate precipitation–dissolution reactions (CaCO₃ + H₂O + CO₂ → Ca²⁺ + 2HCO₃⁻
249 and Ca²⁺ + 2HCO₃⁻ → CaCO₃ + H₂O + CO₂) and mineral carbonation (MgSiO₄ + 2CO₂
250 → 2MgCO₃ + SiO₂ and CaMgSi₂O₆ + CO₂ + H₂O → Ca₂Mg₅Si₈O₂₂(OH)₂ + CaCO₃ +
251 SiO₂) (Mi et al, 2008; Rey, 2015; Yang et al, 2012; Yang and Yang, 2020). These abiotic
252 factors were proved to have large impacts on the dissolution and deposition processes
253 of inorganic C and ultimately determined the reservation and distribution of SIC (Rey,
254 2015; Rowley et al, 2018).

255 However, many biological processes and factors were not quantitatively considered
256 in previous studies. In this study, based on the approach of large-scale field samplings
257 across Tibetan alpine grasslands, we estimated the predominant drivers of SIC stock in
258 the topsoil and subsoil. Our results found the predominant roles of microbial and plant
259 factors in determining SIC stock in both topsoil and subsoil. More importantly, the
260 effects of biotic factors on SIC stock weakened with soil depth (Fig. 4). These results
261 were different from those demonstrating the critical influence of abiotic processes on



262 SIC stock (Mi et al, 2008; Yang et al, 2010).

263 We found that increasing plant aboveground biomass, plant coverage, and root
264 biomass significantly decreased SIC density (Figs. 2 and 3). Plant factors could
265 contribute to the decline of SIC stock by three pathways including uptakes of
266 exchangeable cations, plant organic matter inputs, and rhizosphere processes. First, a
267 large decline in soil base cations is likely to be induced by plant uptake with increasing
268 plant biomass. And the losses of soil exchangeable base cations can cause the
269 transformation of SIC to CO₂, which is ultimately released into the atmosphere (Huang
270 et al, 2015). Second, increasing plant residue inputs can enhance carbonic and organic
271 acid production into soil water solution via microbial decomposition, which reduces the
272 availability of soil base cations through cation exchange in the soil (Sartori et al, 2007)
273 and increase the dissolution and leaching of carbonates, resulting in a decrease in the
274 SIC. Third, the plant rhizosphere effect on releasing CO₂ from carbonates should not
275 be ignored, especially in alkaline soils. By releasing organic acids and protons as well
276 as CO₂, plant roots can reduce soil pH and increase CO₂ in the rhizosphere (Lenzowski
277 et al, 2018), both of which dissolve carbonates by neutralization (Harley & Gilkes,
278 2000). In addition, organic compounds from plant root exudates, such as malate or
279 citrate, can stimulate mineral weathering by dissolving silicate minerals (Dontsova et
280 al, 2020).

281 Furthermore, the topsoil has a larger quantity and higher quality of plant residues
282 than the subsoil, which indicates a more potential for carbonate dissolution by
283 biological processes for the surface soil (Liu et al, 2020). The large root biomass in the



284 topsoil can increase the uptake of base cations and result in increasing proton and
285 organic acids in root exudates (Li et al, 2007), thus reducing the soil carbonate content
286 for maintaining the charge balance. In addition, the larger plant roots exuded more
287 organic compounds in the topsoil that can stimulate parent mineral weathering and
288 dissolve silicate minerals by chelating reaction products (Doetterl et al, 2015; Dontsova
289 et al, 2020).

290 Previous studies reported that microbial properties may not be important in
291 mediating SIC accumulation (Liu et al, 2021; Wang et al, 2015). However, our results
292 found that microbial factors including microbial biomass and bacterial and fungal gene
293 abundance showed significant and negative associations with SIC stock (Figs. 2 and 3),
294 which could be due to microbes driving the carbonate dissolution processes, including
295 microbial respiration, organic matter mineralization, and releases of proton and organic
296 acids by microbial metabolic activity. First, the increase in microbial respiration can
297 improve CO₂ production and enhance the partial pressure of CO₂, leading to a decline
298 in pH and further dissolution of carbonates (Chang et al, 2012). In addition, soil organic
299 matter mineralization and litter decomposition by microbes can induce the dissolution
300 of CO₂ and the release of organic acids (Goulding, 2016; Kuzyakov & Razavi, 2019),
301 both of which decrease the SIC stock. Meanwhile, chelates and enzymes excreted by
302 microbes may contribute to enhancing mineral dissolution rates and organic matter
303 decomposition (Xiao et al, 2015; Zaharescu et al, 2020).

304 We also revealed that bacterial and fungal gene abundance contributed significantly
305 to the variation of SIC stock (Figs. 2 and 3), which was likely to account for decreasing



306 soil pH in the involvement of microbial biological reactions. For instance, nitrifying
307 bacteria can oxidize ammonium to nitrate ($\text{NH}_4^+ + \text{OH}^- + 2\text{O}_2 \rightarrow \text{NO}_3^- + 2\text{H}_2\text{O} + \text{H}^+$),
308 and the production of acidity is finally neutralized through accelerating carbonate
309 dissolution (Zamanian et al, 2016). Also, some nitrogen-fixing bacteria that lived in
310 symbiosis with leguminous plants can acidify the soil by excreting protons during N_2
311 fixation (Vicca et al, 2022). Furthermore, fungi are likely to accelerate carbonate
312 neutralization by exuding protons and organic acids (Van Hees et al, 2006; Wild et al,
313 2021).

314 Microbial factors also affected SIC stock more in the topsoil than in the subsoil.
315 The large plant residues incorporated into the topsoil provided substantial amounts of
316 organic matter for microbial living and decomposition (Oelkers et al, 2015; Ven et al,
317 2020), which can stimulate microbial abundance and activities and promote microbial
318 extracellular enzymes. These extracellular excretions play a fundamental role in
319 microbial respiration and CO_2 production, both of which stimulate silicate weathering
320 and carbonate dissolution (Vicca et al, 2022). Meanwhile, the higher CO_2 flux and CO_2
321 partial pressure resulting from the biological activities of roots and soil microorganisms
322 in the topsoil could enhance carbonate dissolution and formations of pedogenic
323 inorganic C (Chang et al, 2012; Zamanian et al, 2016).

324 Different from plant and microbial factors, the effects of edaphic factors on SIC
325 stock strengthened with soil depth, with soil pH being the most important predictor
326 among edaphic variables (Fig. 4). The buffering capacity in soil solutions determines
327 the equilibrium of ion inputs and outputs by soil pH (Huang et al, 2015). In this study,



328 soil pH in the subsoil (7.85) was much higher than that (7.66) in the topsoil (Table 1).
329 The higher pH could buffer the replacement of the exchangeable cations with protons
330 (Frank & Stuanes, 2003) and increase the preservation of base cations (Gandois et al,
331 2011). Given that base cations and carbonates provide the major buffering capacity in
332 the alkaline soil (Yang et al, 2012), the topsoil could be subject to a larger loss of base
333 cations and SIC due to the lower soil pH compared to the subsoil.

334 Taken together, our results revealed that SIC stock was closely linked with biotic
335 factors, which highlights the roles of biological processes in regulating SIC dynamics
336 (Hong et al, 2019). These results imply that the widespread enhancement of vegetation
337 productivity under global environmental changes (e.g, warming and rewetting) (Ding
338 et al, 2017; Wang et al, 2008) may aggravate the depletion of SIC stock (Raza et al,
339 2020). Meanwhile, previous studies have urged the need for incorporating microbial
340 processes and indicators into Earth system models (ESMs) to reduce the uncertainty in
341 predicting soil C dynamics, especially SOC decomposition (Allison et al, 2010;
342 Moorhead and Sinsabaugh, 2006; Todd-Brown et al, 2013). However, our findings
343 highlighted the vital role of microbial factors in regulating soil C balance from
344 inorganic C preservation. Thus, incorporating microbial processes into the models can
345 aid in the understanding of overall soil C responses, because SOC and SIC are formed,
346 protected, and lost in different ways.

347 More importantly, the effects of biotic factors on SIC stock weakened with soil
348 depth, which implies that SIC may be susceptible to environmental changes in the
349 topsoil where is the hotspot of root and microbial activities. Even though biotic factors



350 in the subsoil played less roles in affecting SIC stock compared with the topsoil, an
351 increase in rooting depth is expected in response to climate warming and land-use
352 change (Liu et al. 2018), which is likely to cause SIC losses in the deep soil by root
353 growth. Therefore, it is a necessity to further explore the effects of biotic factors on SIC
354 stock in the deep soil in the context of global changes. Overall, the contribution of SIC
355 to CO₂ is not ignored and SIC maintenance has a considerable significance on soil C
356 losses and maintains the health and ecosystem functions (Raza et al, 2020; Zamanian
357 et al, 2018). Our study provides robust evidence that biotic factors are mainly
358 responsible for the variation of SIC stock and that topsoils and subsoils should be
359 considered separately when modeling SIC dynamics and its feedbacks on climate
360 change (Yang et al, 2012; Zamanian & Kuzyakov, 2019).

361 **5 Conclusions**

362 Our findings showed that the climatic, edaphic, plant, and microbial variables jointly
363 affected SIC stock in the Tibetan grasslands and that biotic factors had a larger
364 contribution than abiotic factors to the variation of SIC stock. Furthermore, the effects
365 of microbial and plant variables on SIC stock weakened with soil depth, while the
366 effects of edaphic variables strengthened with soil depth. The contrasting responses and
367 drivers of SIC stock between the topsoil and subsoil highlight differential mechanisms
368 underlying SIC preservation with soil depth, which is crucial to understanding and
369 predicting SIC dynamics and its feedbacks to environmental changes.



370 **Data availability.**

371 The data that support the findings of this study are available from the corresponding
372 author upon reasonable request.

373 **Supplement.**

374 Supporting information is also available as supplementary material.

375 **Author contributions.**

376 JP, JW, and SN designed the study. JP, JW, DT, RZ, YL, LS, JY, CW, and SN were
377 involved in drafting or revising the manuscript. All authors read and approved the
378 final manuscript.

379 **Competing interests.**

380 The authors declare that they have no conflict of interest.

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628

Figure captions

629 **Figure 1.** Soil inorganic C content, bulk density, and SIC density in the topsoil and
630 subsoil. The horizontal solid and hollow lines inside each box represent medians and
631 mean values, respectively. Significant differences between the topsoil and subsoil were
632 inspected according to Tukey's test.

633 **Figure 2.** SIC density in relation to climatic, edaphic, plant, and microbial factors in
634 the topsoil. The solid lines are fitted by ordinary least-squares regressions, and the
635 shadow areas correspond to 95% confidence intervals. AM: alpine meadow; AS: alpine
636 steppe; AD: alpine desert; MAP: mean annual precipitation; PAB: plant aboveground
637 biomass; PC: plant coverage. The abbreviations for other variables are shown in Table
638 1. * $p < 0.05$; ** $p < 0.01$; *** $p < 0.001$.

639 **Figure 3.** SIC density in relation to climatic, edaphic, plant, and microbial factors in
640 the subsoil. The solid lines are fitted by ordinary least-squares regressions, and the
641 shadow areas correspond to 95% confidence intervals. AM: alpine meadow; AS: alpine
642 steppe; AD: alpine desert.

643 **Figure 4.** Relative effects of multiple drivers of SIC density in the topsoil (A) and
644 subsoil(B). Climatic variables include MAP, MAT, and altitude; edaphic variables
645 include pH, AN, and sand proportion; plant variables include PB, PC, and RB;
646 microbial variables include MBC, BA, and FA.

647 **Figure 5.** Variation partitioning analyses (VPA) reveal the relative contribution of
648 abiotic and biotic variables to SIC density in the (A) topsoil (61.2% vs. 84.4%) and (B)
649 subsoil (73.4% vs. 86.1%), respectively. Results in three fractions: the unique effect of

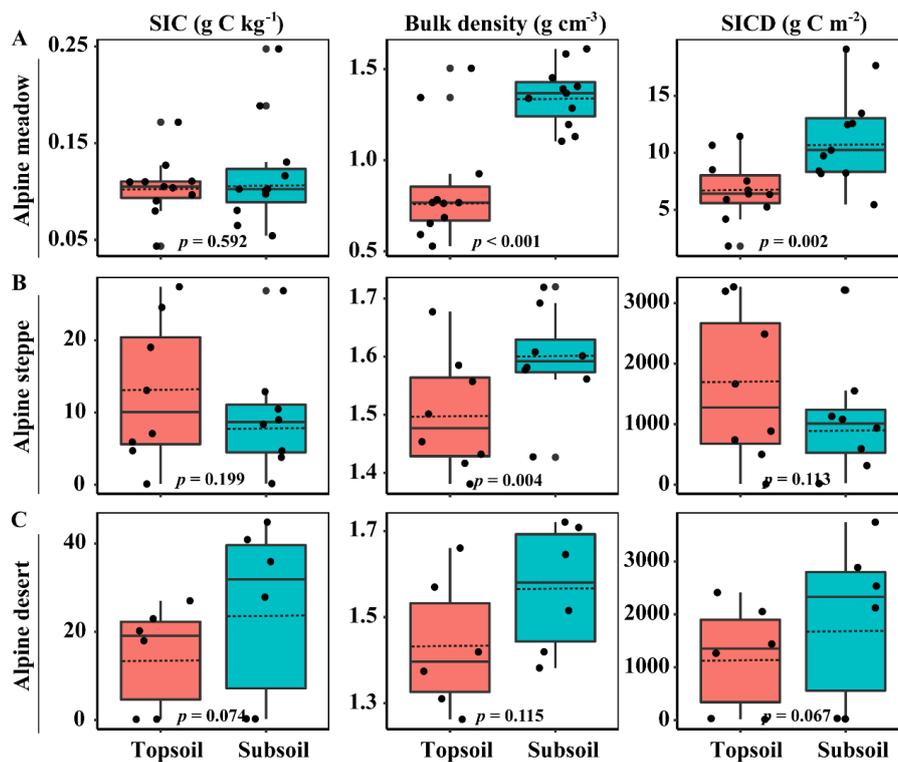


650 abiotic factors (X1), the unique effect of biotic factors (X2), and common interception

651 of abiotic and biotic factors (X3).



652 **Figure 1.**
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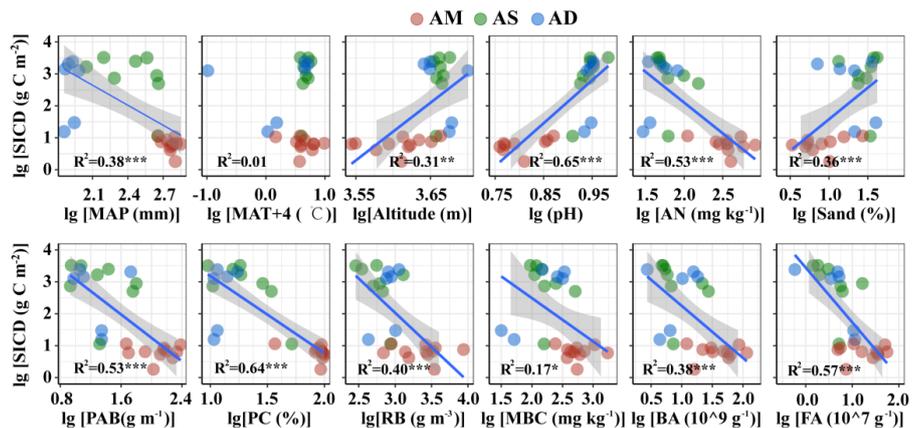


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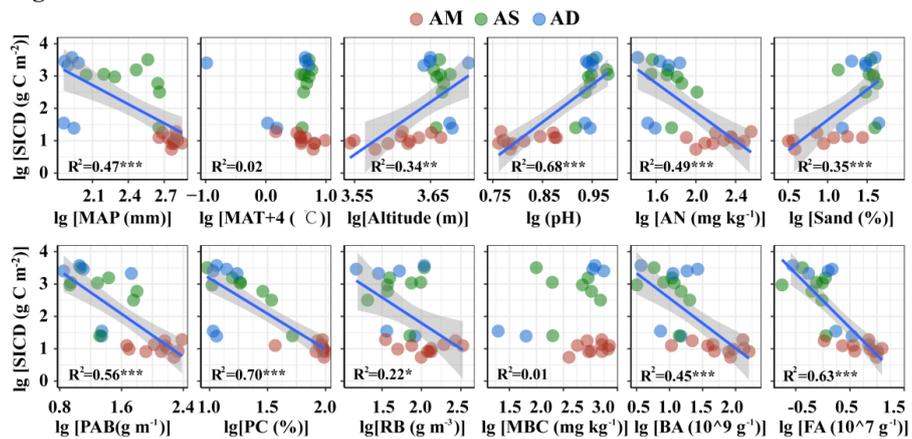
656 **Figure 2.**



657



658 **Figure 3.**

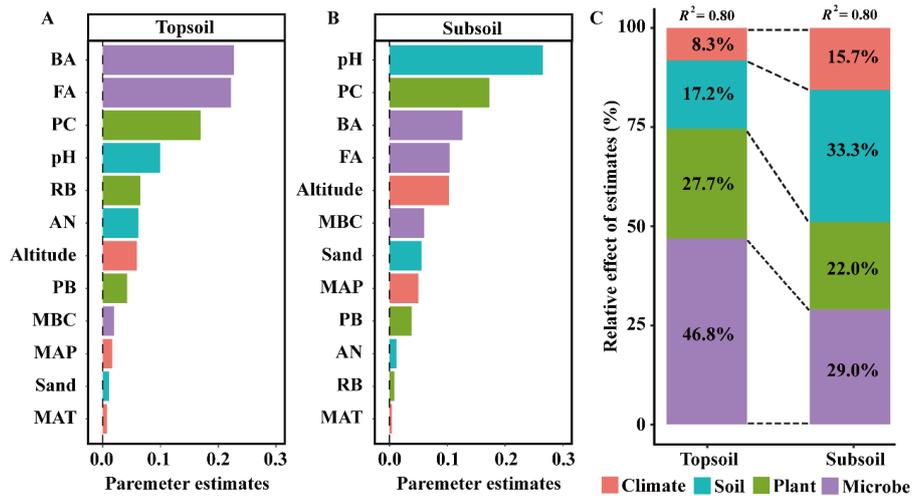


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661 **Figure 4.**

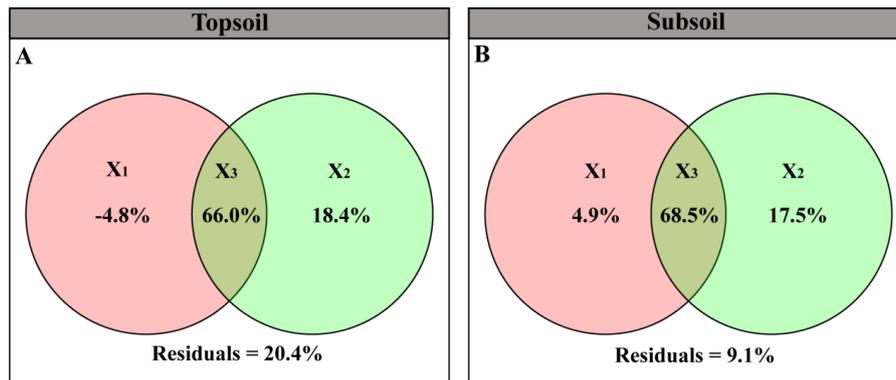


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664 **Figure 5.**



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667 **Table 1.** Edaphic, plant, and microbial properties between the topsoil and subsoil for
668 25 sampling sites.

Parameters	Topsoil	Subsoil	<i>p</i> value
RB (g m ⁻²)	1670 ± 359	95.2 ± 15.3	<0.001
pH	7.66 ± 0.28	7.85 ± 0.26	<0.001
AN (mg kg ⁻¹)	217 ± 43.7	131 ± 22.0	0.004
SP (%)	47.1 ± 4.33	45.6 ± 4.87	0.698
MBC (mg kg ⁻¹)	385 ± 73.8	101 ± 9.7	0.001
BA (10 ⁹ gene copies g ⁻¹ soil)	27.2 ± 5.68	12.6 ± 2.86	0.001
FA (10 ⁷ gene copies g ⁻¹ soil)	14.2 ± 3.25	3.62 ± 0.84	0.001

669 RB: root biomass; AN: soil available nitrogen; SP: sand proportion; MBC: microbial
670 biomass carbon; BA: soil bacterial abundance; FA: soil fungal abundance. Values are
671 means ± standard error (SE). *p* values represent significant differences between the
672 topsoil and subsoil according to Tukey's test.



673 **Supporting information**

674 Additional supporting information may be found online in the supporting information

675 tab for this article.