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

2 **Tibetan alpine grasslands**

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Abstract. The soil inorganic carbon (SIC) pool is a major component of soil C pools,

and clarifying the predictors of SIC stock is urgent for decreasing soil C losses and

maintaining soil health and ecosystem functions. However, the drivers and their relative

effects on the SIC stock at different soil depths remain largely unexplored. Here, we

conducted a large-scale sampling to investigate the effects and relative contributions of

abiotic (climate and soil) and biotic (plant and microbe) drivers on the SIC stock

between topsoils (0–10 cm) and subsoils (20–30 cm) across Tibetan alpine grasslands.

Results showed that the SIC stock had no significant differences between the topsoil

and subsoil. The SIC stock showed a significant increase with altitude, pH, and sand

proportion, but declined with mean annual precipitation, plant aboveground biomass,

plant coverage, root biomass, soil available nitrogen, microbial biomass carbon, and

bacterial and fungal gene abundance. For both soil layers, biotic factors had larger

effects on the SIC stock than abiotic factors did. But the relative importance of these

determinants varied with soil depth, with the effects of plant and microbial variables on

SIC stock weakening with soil depth, whereas the importance of climatic and edaphic

variables increasing with soil depth. Specifically, bacterial and fungal gene abundance

and plant coverage played dominant roles in regulating SIC stock in the topsoil, while

soil pH contributed largely to the variation of SIC stock in the subsoil. Our findings

highlight differential drivers over SIC stock with soil depth, which should be considered

in biogeochemical models for better simulating and predicting SIC dynamics and its

feedbacks to environmental changes.

1 Introduction

Soils store approximately 1,500 Pg of organic carbon (SOC) and 940 Pg of inorganic carbon (SIC) to a depth of 1 m (Batjes, 1996; Jobbágy & Jackson, 2000), which are the largest carbon (C) pool in the terrestrial ecosystem and play a critical part in the global C cycling (Darwish et al., 2018; Lal 2004; Prietzel et al, 2016). To alleviate the elevated levels of atmospheric carbon dioxide, most previous studies concentrate on the SOC pool because it responds quickly to global climate change such as warming and nitrogen deposition, and it is strongly linked with various ecosystem functions (Wang et al., 2002; Yang et al., 2012). Compared to the relatively short turnover time of SOC, SIC has a long residence time due to soil weathering (Monger et al, 2015; Zang et al, 2018), which is considered to be fairly stable and has less contribution to changes in terrestrial ecosystem C balance (Yang et al, 2012). Therefore, previous studies have paid little attention to SIC. However, recent studies suggest that SIC is also responsive to anthropogenic activities and global climate changes such as soil acidification, atmospheric N deposition, and global warming (Yang et al, 2010; Song et al, 2022), acting as a critical C source (Liu et al, 2020) or C sink (Gao et al, 2018; Liu et al, 2021). Thus, the preservation of SIC and its roles in climate mitigation should not be neglected, especially in arid and semi-arid grasslands which store a large amount of SIC (Yang et al, 2012).

SIC stock and stability can be fundamentally altered by an array of abiotic and biotic processes (Raza et al, 2020). High precipitation can promote soil silicate minerals weathering and removal of base cations (Ca^{2+}, Mg^{2+}, K^+, and Na^+) by leaching (Vicca

et al, 2022). Soil acidification due to atmospheric nitrogen (N) and acid deposition and the nitrification of NH_4^+ may greatly accelerate soil carbonate dissolution and CO_2 releases (Raza et al, 2020; Song et al, 2022). Plant growth can deplete soil carbonates by releasing proton and organic acids from root rhizosphere (Goulding et al, 2016; Kuzyakov & Razavi, 2019), and biological N_2 fixation by some legumes are likely to cause SIC losses (Tang et al, 1999). Furthermore, plant autotrophic and microbial heterotrophic respiration often facilitate carbonate dissolution by enhancing CO_2 partial pressures (An et al, 2019; Liu et al, 2021). Nevertheless, how these abiotic and biotic factors affect SIC stock and what is the relative importance of these confounding drivers remain largely uncertain.

Previous studies on SIC stock mostly have focused on the topsoil within 10 cm soil depth (Yost and Hartemink, 2020), which have relatively different from the subsoil (i.e., soils residing >20 cm below ground) in the aspect of biochemical processes, plant roots, soil properties, and microbial communities (Rumpel et al., 2012; Zhou et al., 2021), while the patterns of SIC stock in the subsoil on a large scale remain elusive. The predictors of SIC stock in the subsoil may differ from those in the topsoil due to distinct soil microenvironments, soil physicochemical properties, root exudates, and microbial abundance and functions (Jia et al, 2017). For instance, the topsoil has larger root biomass and higher microbial activity than the subsoil, but the subsoil tends to preserve soil parent material because of the weakened weathering by the isolation of heat and energy from the surface soil (Crowther et al, 2016). Thus, the abiotic and biotic

variables may exhibit different effects on SIC stock in the subsoil compared to the topsoil due to the various importance of these variables.

The Tibetan Plateau has the largest alpine grassland on the Eurasian continent, which is a vital component of global terrestrial ecosystems, providing an ideal platform to explore SIC stock and its determinants (Wang et al, 2002; Yang et al, 2010). During the past several decades, the plateau has experienced significant warming (Wang et al, 2008) and pronounced atmospheric N deposition (Liu et al, 2013; Yu et al, 2019). This continuous warming and N deposition have resulted in a significant increase in plant growth and soil acidification (Ding et al, 2017; Yang et al, 2012), which could be likely to induce potential CO_2 releases from soil carbonates by biogeochemical process (Raza et al, 2020). However, a general understanding of SIC stock with soil depth across Tibetan alpine grasslands remains unexplored.

Here, we researched the relative importance of climatic, edaphic, plant, and microbial variables to SIC stock at different soil layers along an approximately 3,000 km transect of alpine grasslands on the Tibetan Plateau, spanning a broad range of climatic and geographical conditions. Specifically, two key questions are addressed in this study: (1) what are the differences of SIC stock between the topsoil and subsoil? (2) how does the relative importance of climatic, edaphic, plant, and microbial variables to the variation of SIC stock along with soil depth?

2 Materials and methods

2.1 Study area and field sampling

From30 July to 28 August 2020, we conducted large-scale systematic field surveys and samplings in Tibetan alpine grasslands. The total 25 sampling sites covered approximately 3,000 km and included three grassland types (i.e, 11 alpine meadow, 8 alpine steppe, and 6 alpine desert sites). The distance between nearby sampling sites was about 120 km. The study sites cover a broad geographic and climatic range, with longitude and latitude ranging from 79°49'39" to 102°25'31" E and 31°06'37" to 32°43'09" N, respectively, and the altitude ranging from 3500 m to 5016 m. These sites covered a broad precipitation gradient varying between 72 mm and 706 mm. The mean annual temperature (MAT) ranged from –3.9°C to 5.8°C. The plant communities were dominated by *Kobresia tibetica Maxim*, *Stipa caucasica*, *Kobresia pygmaea*, *Stipa purpurea*, and *Leontopodium pusillum*. Soils were *Cambisol* and some were loess-derived *Luvisol*. The site location, grassland type, climatic, and plant parameters were detailed in Table S1.

2.2 Climatic data

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

2.3 Soil properties

At each site, we selected four 1 m ×1 m plots for soil and plant samplings and the distance between nearby sampling plots was 25 m. In each plot, a 7.5-cm diameter soil drill was used to take five soil cores at fixed soil depths (0-10 cm, 10-20 cm, and 20-30 cm), and a 2-mm mesh was used to remove stones. Based on our field observation, the soil depth is relatively shallow (less than 40 cm) for alpine grasslands, especially for the alpine desert. Moreover, most of the belowground roots in alpine grasslands distribute on the surface of 10 cm and decrease sharply below 20 cm. Thus, we defined the topsoil and subsoil as 0-10 cm and 20-30 cm soils, respectively. After mixing, 100 g of fresh soils from each plot were collected and stored in a –4°C portable icebox, then returned to the laboratory and stored at –20°C for microbial properties. The rest soil samples, about 700 g, were also sent back to the laboratory and air-dried for measurements of other soil properties including soil pH, AN, and mechanical composition. A 40 cm ×40 cm ×40 cm (length × width × depth) pit was dug for measuring soil bulk density (BD) by using a constant volume soil sampling drill (100 cm^3), and the undisturbed soil was preserved in aluminum specimen boxes returning to the laboratory and oven-dried for 48 hours at 105°C and weighed. The oven-dried soil (20 g) was screened for gravel by sifting through a 2-mm mesh sieve and material larger than 2 mm were collected and weighed to determine the percentage of gravels. Soil pH (1:25 soil: H_2O) was measured using a soil pH meter, and available nitrogen (AN) was determined by the alkaline-hydrolysis diffusion method. A laser particle analyzer (Mastersizer 2000, Malvern Panalytical, UK) was applied to measure soil mechanical

142   compositions, including clay (< 2 μm), silt (2-50 μm), and sand (> 50 μm) proportion.

143   SIC was determined by using an inorganic C analyzer (multi EA® 4000; Analytic Jena,

144   Germany). The multi EA 4000 C elemental analyzer was equipped with the automatic

145   TIC solids module and calibrated before the analysis. The sample boat was acidified

146   automatically with 40 % H_3PO_4 in the reactor of the TIC module. And the CO_2 from

147   the carbonate was released, the measuring gas was dried and cleaned and the carbon

148   content was measured by means of the wide-range NDIR detector. Before being

149   analyzed directly, all soil samples were ground into solid fine powders with a mortar,

150   and for the determination of TIC, a standard, prepared by solids-dilution of $CaCO_3$ with

151   SiO_2 (0.2 % C), was used, with weighting rage 7-200 mg, to cover a wide concentration

152   range.

153   **2.4 Plant properties**

154   In each plot, we estimated plant coverage (PC) by the projection method, namely the

155   proportion of vegetation projection to the area of the sampling plot. In addition, plant

156   aboveground biomass were clipped to ground level and collected, belowground roots

157   were sampled by 3 soil samples in each plot, which was mixed by 2 soil cores with 7.5

158   cm diameter drill, and collected from soil by rinsing in water, Finally, they were oven-

159   dried at 60°C and weighed to determine plant aboveground biomass (PAB) and root

160   biomass (RB) , respectively.

161   **2.5 Microbial attributes**

Soil microbial biomass carbon (MBC) was measured by using a chloroform fumigation-extraction procedure (Brookes et al, 1985). Briefly, 10 g of unfumigated and chloroform-fumigated fresh soil samples were extracted by using 0.5 M K_2SO_4 after 24 h of incubation, respectively. Then, the extracts were analyzed by using a TOC analyzer (multi N/C® 3100; Analytic Jena, Germany). The MBC was determined by the differences in C concentrations between unfumigated and chloroform-fumigated samples, and the correction factor (i.e, KC= 0.45) was used to convert microbial C to MBC (Joergensen, 1996).

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

2.6 Statistical analyses

The total SIC density (C stock per land area) in each soil depth layer was calculated

using Equation (1) (Pan et al, 2019):

SIC density (g C m^{-2}) = SIC (g C kg^{-1}) × BD (g cm^{-3}) ×d (cm) × (1-g) /100   (1)

where SIC is soil inorganic C content, d is the depth of the soil layer (0.1 m), BD is

bulk density, and g is the percentage of gravel fraction (>2 mm).

First, the differences of SIC stock and corresponding abiotic and biotic variables

between the topsoil and subsoil were examined by *T*-test. Second, SIC density and

various abiotic and biotic variables were log-transformed and standardized (z-score

normalization) to perform the assumption of normality and homogeneity by Shapiro-

Wilk and Levene's test, respectively (Pan et al, 2021). Then the linear regressions were

used to test SIC density about different variables for both the topsoil and subsoil across

sites. Also, the Pearson correlation coefficients between SIC density and each variable

were analyzed in Table S3.

Third, a linear model was employed to examine SIC density with abiotic and biotic

variables by using the maximum likelihood estimation with the lm package. And the

relative effect of all parameter estimates was calculated to evaluate the relative

importance of drivers SIC density. Each predictor variable was simultaneously tested

in the model, which was comparable for the contribution of different types of predictor

factors to SIC density. And the absolute values of standardized regression coefficients

of the explanatory variables accounting for the percentage of the sum of all standardized

regression coefficients were used to express the importance of predictors (Gross et al.,

2017; Le Provost et al., 2020). Also, SIC density and abiotic and biotic variables were

206 standardized before analyses, using the Z-score to interpret variable estimates on a

207 comparable scale (Gross et al, 2017).

208 Log (SIC density) = $\beta_0 + \beta_1 \log X_1 + \beta_1 \log X_2 + \ldots \beta_{12} \log X_{12}$ (2)

209 where β_0 and β_i (i=1, 2, 3…12) are intercept and coefficients, respectively. To

210 explore the predictors of SIC density in different soil depths across all sites, the absolute

211 values of slopes of the variables were extracted and plotted. Then, to quantify their

212 relative contribution to SIC density, 12 predictor variables were categorized into four

213 groups, including climatic (MAP, MAT, and altitude), edaphic (pH, AN, and sand

214 proportion), plant (PB, PC, and RB), and microbial (MBC, BA, and FA) factors, the

215 detailed categorization of explanatory variables was listed in Table S2.

216 Furthermore, the relative importance of abiotic (climatic and edaphic) and biotic

217 (plant and microbial) variables in predicting SIC density was quantified by performing

218 variation partitioning analyses (VPA) (Borcard et al., 1992) and using the "vegan"

219 package in R 4.1.3, which was used to divide the variation of SIC density among two

220 types of explanatory variables for their individual and joint effects In this analysis, the

221 common and unique contribution of sets of explanatory variables (two sets including

222 abiotic and biotic variables) in SIC density is determined. And, the residuals were

223 determined by a fraction of response variables variations, which could not be explained

224 by any of the explanatory variables. The VPA method allows us to explore the variation

225 clearly by the percentage of explanatory variables, which are easy to interpret and can

226 be discussed in the context of SIC density.

227

3 Results

3.1 SIC density and influencing variables in different soil depths

SIC density and SIC content had no significant differences between the topsoil and subsoil, but bulk density in the subsoil was much higher compared with the topsoil. Specifically, SIC density in the topsoil and subsoil ranged from 1.8 g C m^{-2} to 3271 g 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 ± 220 g C m^{-2} and 814 ± 236 g C m^{-2}, respectively (Fig. 1). No significant changes in SIC density with soil depth were observed in both the alpine steppe and alpine desert (p=0.113 and p=0.068, respectively; Fig. 1), but SIC density was higher in the subsoil than that in the topsoil in the alpine meadow ($p = 0.002$, Fig. 1).

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

3.2 Associations of SIC density with abiotic and biotic variables

The SIC density was closely related to multiple abiotic and biotic variables (Table S3, Figs. 2 and 3 for topsoil and subsoil, respectively). For both the topsoil and subsoil, the SIC density showed a significant increase trend with altitude, pH, and sand proportion, but declined with MAP, PAB, PC, RB, AN, BA, and FA (all p<0.05). The SIC density

showed a correlation with MBC in the topsoil ($p<0.05$, Fig. 2), but not in the subsoil (Fig. 3). Meanwhile, the SIC density in both two soil depths did not significantly correlate with MAT (Figs. 2 and 3). Also, the absolute value of slope for the regression equation for the most explanatory variables (except for AN, MAT, and MBC) in the topsoil was larger than that of the subsoil, especially for RB and SP (Figs. 2 and 3).

3.3 Determinants of SIC density in different soil depths

The linear model and VPA collectively displayed that the predominant predictors of SIC density differed with soil depth (Figs. 4 and 5). Specifically, for the topsoil, the linear model revealed that microbial and plant variables largely explained the variations in the SIC density, followed by edaphic variables and climate contributed the least (Fig. 4). Among these variables, PC, BA, and FA exhibited larger effects on the SIC density compared with other predictor factors (Fig. 4). Also, the VPA analysis illustrated that biotic factors explained the majority variation of SIC density compared with abiotic factors (Fig. 5). For the subsoil, the linear model showed that edaphic variables largely explained the variation in SIC density, followed by microbial and plant variables, and climate contributed the least (Fig. 4). Among these variables, the soil pH had larger contributions to the variation of SIC density rather than others (Fig. 4). Meanwhile, the VPA analysis confirmed that the predict of biotic factors on SIC density were better than those of abiotic factors in the subsoil (Fig. 5).

4 Discussion

To the best of our knowledge, this study was the first to study large-scale evidence of the relative contribution of abiotic and biotic drivers to the variation of SIC stock at different soil depths, which has considerable implications for grasping the importance of SIC in the ecosystem C cycling. Due to considerably stable characteristics and the long turnover time (Mi et al, 2008; Yang et al, 2010; Zamanian et al, 2018), SIC stock is traditionally considered to be dominated by abiotic factors including soil moisture, soil pH, CO_2 partial pressure, and Ca^{2+} concentrations according to the equilibrium of carbonate precipitation–dissolution reactions ($CaCO_3 + H_2O + CO_2 \rightarrow Ca^{2+} + 2HCO_3^-$ and $Ca^{2+} + 2HCO_3^- \rightarrow CaCO_3 + H_2O + CO_2$) and mineral carbonation ($MgSiO_4 + 2CO_2 \rightarrow 2MgCO_3 + SiO_2$ and $CaMgSi_2O_6 + CO_2 + H_2O \rightarrow Ca_2Mg_5Si_8O_{22}(OH)_2 + CaCO_3 + SiO_2$) (Mi et al, 2008; Rey, 2015; Yang et al, 2012; Yang and Yang, 2020). These abiotic factors were proved to have large impacts on the dissolution and deposition processes of inorganic C and ultimately determined the reservation and distribution of SIC (Rey, 2015; Rowley et al, 2018).

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

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

291   We found that SIC density showed a declining trend with the increasing plant

292   aboveground biomass, plant coverage, and root biomass (Figs. 2 and 3). Plant factors

293   could contribute to the decline of SIC stock by three pathways including uptakes of

294   exchangeable cations, plant organic matter inputs, and rhizosphere processes. First, a

295   large decline in soil base cations is likely to be induced by plant uptake with increasing

296   plant biomass. And the losses of soil exchangeable base cations can cause the

297   transformation of SIC to CO_2, which is ultimately released into the atmosphere (Huang

298   et al, 2015). Second, increasing plant residue inputs can enhance carbonic and organic

299   acid production into soil water solution via microbial decomposition, which reduces the

300   availability of soil base cations through cation exchange in the soil (Sartori et al, 2007)

301   and increase the dissolution and leaching of carbonates, resulting in a decrease in the

302   SIC. Third, the plant rhizosphere effect on releasing CO_2 from carbonates should not

303   be ignored, especially in alkaline soils. By releasing organic acids and protons as well

304   as CO_2, plant roots can reduce soil pH and increase CO_2 in the rhizosphere (Lenzewski

305   et al, 2018), both of which dissolve carbonates by neutralization (Harley & Gilkes,

306   2000). In addition, organic compounds from plant root exudates, such as malate or

307   citrate, can stimulate mineral weathering by dissolving silicate minerals (Dontsova et

308   al, 2020).

309   Furthermore, the topsoil has a larger quantity and higher quality of plant residues

310   than the subsoil, which indicates a more potential for carbonate dissolution by

311   biological processes for the surface soil (Liu et al, 2020). The large root biomass in the

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

Also, the SIC density in both two soil depths appears to have an increase or decrease trend from the alpine meadow to the alpine steppe and alpine desert (Figs. 2 and 3). In the present study, for example, the alpine meadow has larger plant productivity than the alpine steppe, which implies that more plant above- and below-ground residues are deposited in alpine meadow soils compared to alpine steppe soils. Therefore, from the perspective of the whole ecosystem, the grassland type would be a better predictor for the quantity and distribution of SIC density.

Previous studies reported that microbial properties may not be important in mediating SIC accumulation (Liu et al, 2021; Wang et al, 2015). However, our results found that microbial factors including microbial biomass and bacterial and fungal gene abundance showed significant associations with SIC stock (Figs. 2 and 3), which could be due to microbes driving the carbonate dissolution processes, including microbial respiration, organic matter mineralization, and releases of proton and organic acids by microbial metabolic activity. First, the increase in microbial respiration can improve CO_2 production and enhance the partial pressure of CO_2, leading to a decline in pH and further dissolution of carbonates (Chang et al, 2012). In addition, soil organic matter

mineralization and litter decomposition by microbes can induce the dissolution of CO_2 and the release of organic acids (Goulding, 2016; Kuzyakov & Razavi, 2019), both of which decrease the SIC stock. Meanwhile, chelates and enzymes excreted by microbes may contribute to enhancing mineral dissolution rates and organic matter decomposition (Xiao et al, 2015; Zaharescu et al, 2020).

We also revealed that bacterial and fungal gene abundance were significantly correlated with SIC stock (Figs. 2 and 3), which was likely to account for decreasing soil pH in the involvement of microbial biological reactions. For instance, nitrifying bacteria can oxidize ammonium to nitrate ($NH_4^+ + OH^- + 2O_2 \rightarrow NO_3^- + 2H_2O + H^+$), and the increase in acidity is finally neutralized through accelerating carbonate dissolution (Zamanian et al, 2016). Also, some nitrogen-fixing bacteria that lived in symbiosis with leguminous plants can acidify the soil by excreting protons during N_2 fixation (Vicca et al, 2022). Furthermore, fungi are likely to accelerate carbonate neutralization by exuding protons and organic acids (Van Hees et al, 2006; Wild et al, 2021).

Microbial factors also could the better predictors for SIC stock in the topsoil than in the subsoil. The large plant residues incorporated into the topsoil provided substantial amounts of organic matter for microbial living and decomposition (Oelkers et al, 2015; Ven et al, 2020), which can stimulate microbial abundance and activities and promote microbial extracellular enzymes. These extracellular excretions play a fundamental role in microbial respiration and CO_2 production, both of which stimulate silicate weathering and carbonate dissolution (Vicca et al, 2022). Meanwhile, the higher CO_2

flux and CO_2 partial pressure resulting from the biological activities of roots and soil microorganisms in the topsoil could enhance carbonate dissolution and formations of pedogenic inorganic C (Chang et al, 2012; Zamanian et al, 2016).

Different from plant and microbial factors, the prediction of edaphic factors on SIC stock strengthened with soil depth, with soil pH being the most important predictor among edaphic variables (Fig. 4). The buffering capacity in soil solutions determines the equilibrium of ion inputs and outputs by soil pH (Huang et al, 2015). In this study, soil pH in the subsoil (7.85) was much higher than that (7.66) in the topsoil (Table 1). The higher pH could buffer the replacement of the exchangeable cations with protons (Frank & Stuanes, 2003) and increase the preservation of base cations (Gandois et al, 2011). Given that base cations and carbonates provide the major buffering capacity in the alkaline soil (Yang et al, 2012), the topsoil could be subject to a larger loss of base cations and SIC due to the lower soil pH compared to the subsoil.

Taken together, our results revealed that SIC stock was closely linked with biotic factors, which highlights the roles of biological processes in predicting SIC dynamics (Hong et al, 2019). These results imply that the widespread enhancement of vegetation productivity under global environmental changes (e.g, warming and rewetting) (Ding et al, 2017; Wang et al, 2008) may aggravate the depletion of SIC stock (Raza et al, 2020). Meanwhile, previous studies have urged the need for incorporating microbial processes and indicators into Earth system models (ESMs) to reduce the uncertainty in predicting soil C dynamics, especially SOC decomposition (Allison et al, 2010; Moorhead and Sinsabaugh, 2006; Todd-Brown et al, 2013). However, our findings

highlighted the vital role of microbial factors in regulating soil C balance from inorganic C preservation. Thus, incorporating microbial processes into the models can aid in the understanding of overall soil C responses, because SOC and SIC are formed, protected, and lost in different ways.

More importantly, the predictions of biotic factors on SIC stock weakened with soil depth, which implies that SIC may be susceptive to environmental changes in the topsoil where is the hotspot of root and microbial activities. Even though biotic factors in the subsoil played a less role in predicting SIC stock compared with the topsoil, an increase in rooting depth is expected in response to climate warming and land-use change (Liu et al. 2018), which is likely to cause SIC losses in the deep soil by root growth. Therefore, it is a necessity to further explore the effects of biotic factors on SIC stock in the deep soil in the context of global changes. Although most of the variations in SIC density were explained by our measured explanatory variables, some other potential variables may also predict SIC density (Fig. 5). Then, understanding the effects of other potential abiotic and biotic factors on SIC density with soil depth is urgently needed when predicting the response and feedback of SIC to climate change in the future. Overall, the contribution of SIC to CO_2 is not ignored and SIC maintenance has a considerable effect on soil C losses and is important to maintain the health and ecosystem functions (Raza et al, 2020; Zamanian et al, 2018). Our study provides robust evidence that biotic factors are correlated with SIC stock in the Tibetan plateau and that topsoils and subsoils should be considered separately when modeling SIC dynamics and its feedbacks on climate change (Yang et al, 2012; Zamanian &

400   Kuzyakov, 2019).

5 Conclusions

402   Our findings showed that SIC stock had no significant differences between the topsoil

403   and subsoil in the Tibetan grasslands and the climatic, edaphic, plant, and microbial

404   variables jointly predicted SIC stock in the Tibetan grasslands, and that biotic factors

405   had a larger contribution than abiotic factors to the variation of SIC stock. Furthermore,

406   the relative importance of explanatory variables to the variation of SIC stock varied

407   with soil depth, the predictions of microbial and plant variables on SIC stock weakened

408   with soil depth, while the predictions of edaphic variables strengthened with soil depth.

409   Our results revealed that biotic factors should be considered seriously for predicting

410   SIC stock due to their regulating roles in biological processes. The contrasting

411   responses and drivers of SIC stock between the topsoil and subsoil highlight differential

412   mechanisms underlying SIC preservation with soil depth, which is crucial to

413   understanding and predicting SIC dynamics and its feedbacks to environmental changes.

Data availability.

The data that support the findings of this study are available from the corresponding

author upon reasonable request.

Supplement.

Supporting information is also available as supplementary material.

Author contributions.

JP, JW, and SN designed the study. JP, JW, DT, RZ, YL, LS, JY, CW, and SN were

involved in drafting or revising the manuscript. All authors read and approved the

final manuscript.

Competing interests.

The authors declare that they have no conflict of interest.

Acknowledgments

This study was financially supported by the Second Tibetan Plateau Scientific

Expedition and Research (STEP) program (2019QZKK0302), the National Natural

Science Foundation of China (31988102 and 32101390), and the China National

Postdoctoral Program for Innovative Talents (BX20200330).

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 Figure captions

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

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

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

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

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

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

702   of abiotic and biotic factors (X3).

Figure 1.

707   **Figure 2.**

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709   **Figure 3.**

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Figure 4.

715 **Figure 5.**

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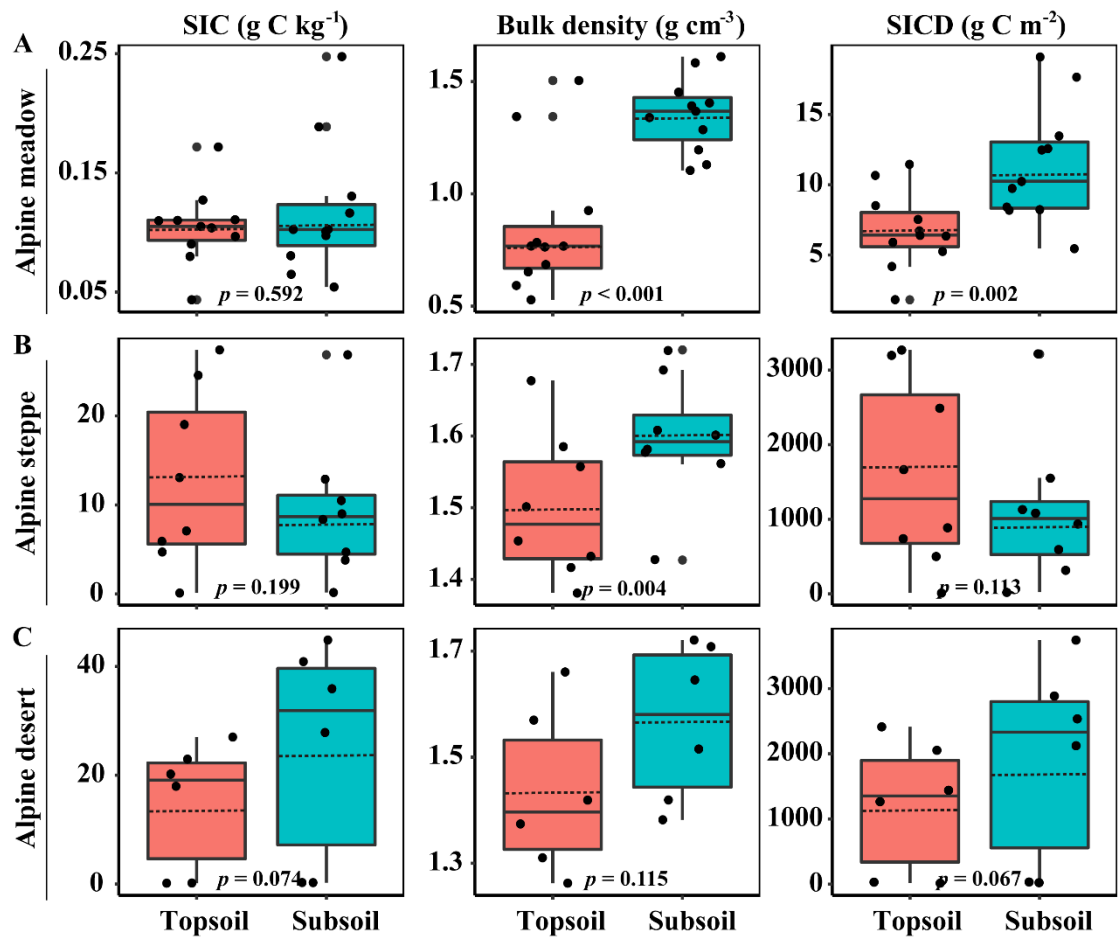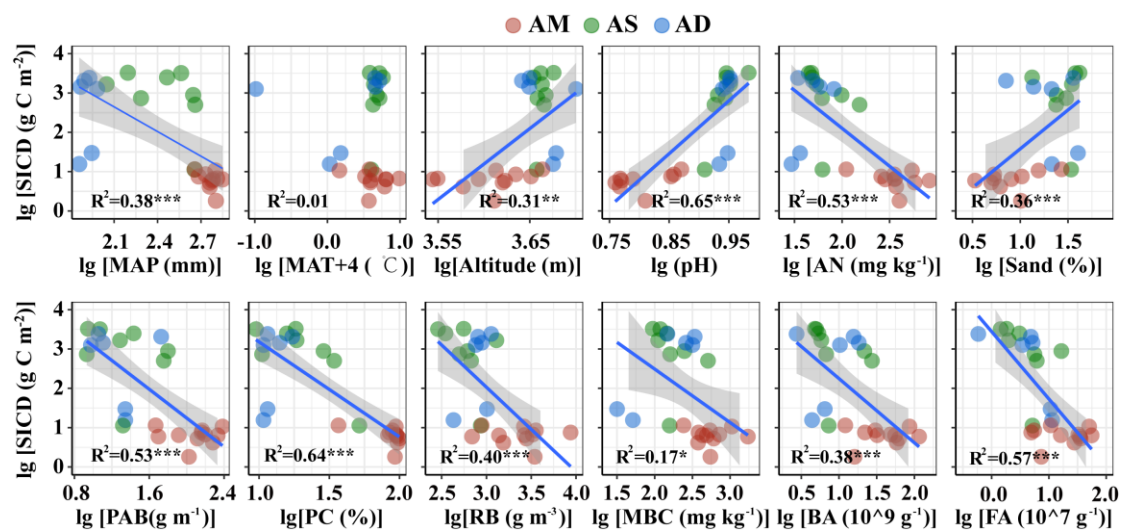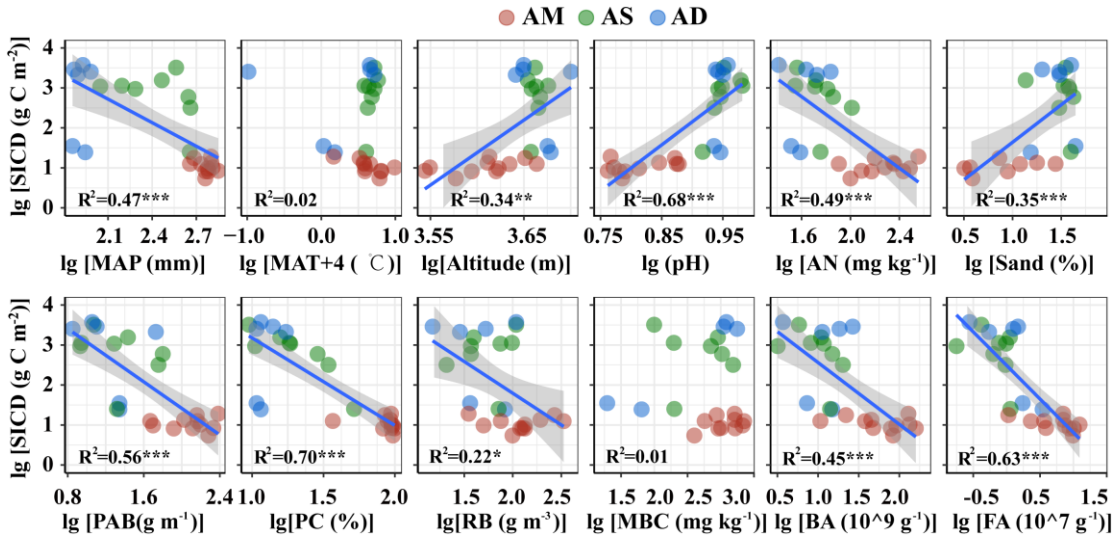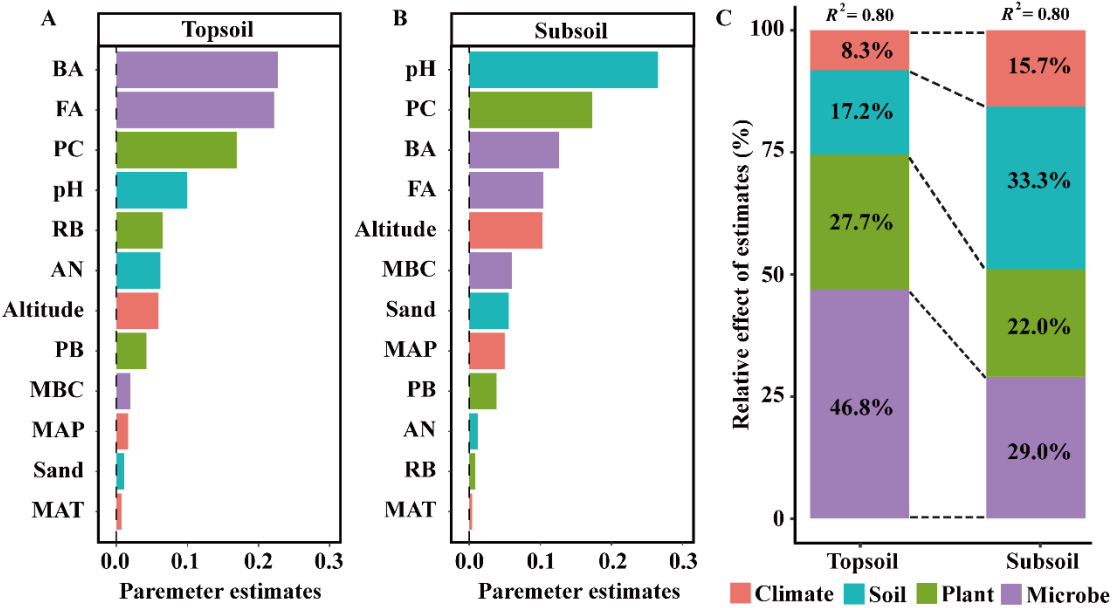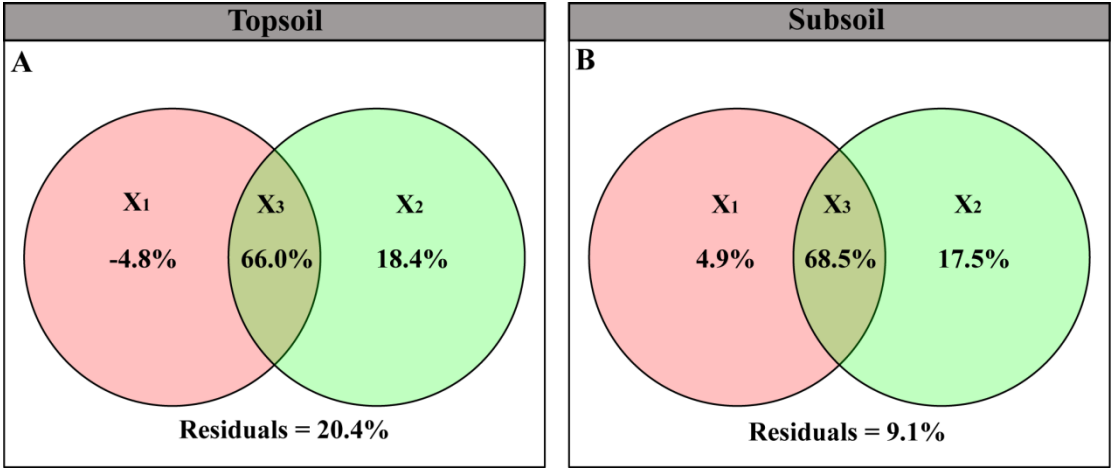
718   **Table 1.** Edaphic, plant, and microbial properties between the topsoil and subsoil for
719   25 sampling sites.

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

720   RB: root biomass; AN: soil available nitrogen; SP: sand proportion; MBC: microbial
721   biomass carbon; BA: soil bacterial abundance; FA: soil fungal abundance. Values are
722   means $\pm$ standard error (SE). *p* values represent significant differences between the
723   topsoil and subsoil according to Tukey's test.

724 **Supporting information**

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

726 tab for this article.