data homogenization processing summary and QC check results: $ChronicN_stocks$ notes included with key file:

source	Var_long	var	var_notes
location	Google Directory		ChronicN_stocks
location	Network (e.g. LTER, CZO, DIRT,	network	LTER
	NutNet, etc)		
location	Site code (e.g. LUQ) or name	$\operatorname{site_code}$	HFR
location	Location name	location_name	Harvard Forest
location	control samples identifier	$control_id$	Trt = 0
location	number of treatments	$number_treatments$	high & low N

files processed:

type	filename
provided data	Harvard Forest_Chronic N Amendment
	Study_Soil C_updated
homogenized data	Harvard Forest_Chronic N Amendment
	$Study_Soil~C_updated_HMGZD$

QC results: location data

location data checks passed

QC results: profile data, data range

var	min	max	minValue	$\max Value$	error
layer_bot	-1	50	0	500	out of range

QC results: profile data, data type

profile data type checks passed