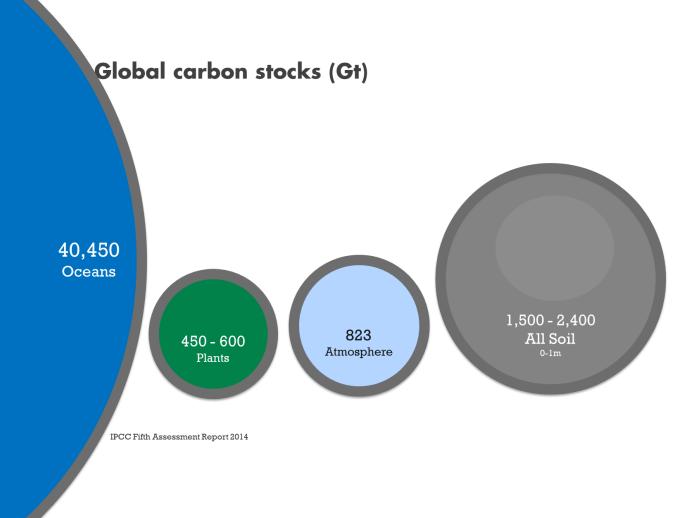


Nature Based Solutions

Christian Davies

Principal Science Expert

Shell International Exploration and Production Inc.





Different types of NBS

NBS comprise all activities related to the protection, or re-development, of natural ecosystems – such as forests, grasslands, wetlands and blue carbon – to lower concentrations of greenhouse gases in the atmosphere.



- Coastal restoration and protection
- Peat restoration and protection



- Avoiding deforestation
- Reforing degraded forest
- Improving forest management
- Afforestaion
- Reforestation

AGRICULTURE & GRASSLANDS

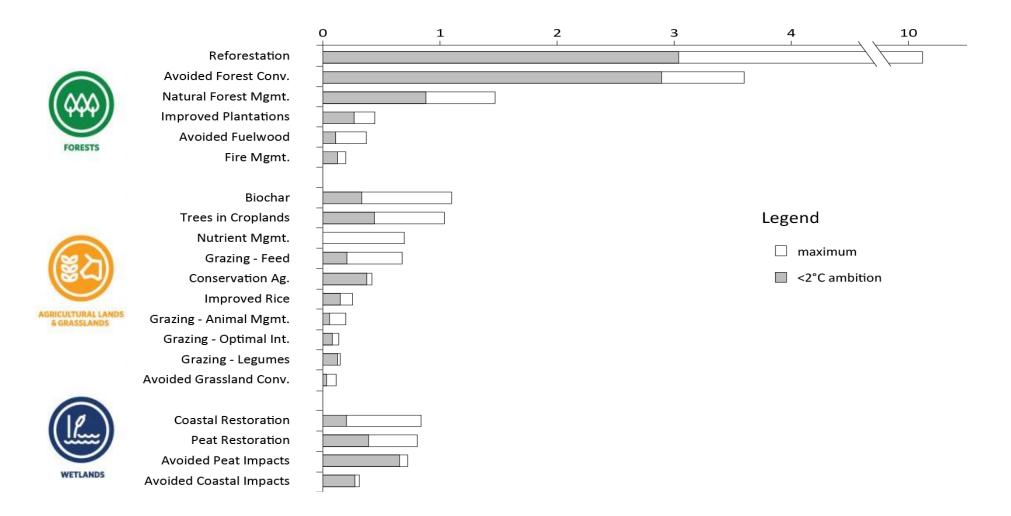
- Biochar for soil amendment
- Restor trees in croplands
- Improve nutrient management
- Improve rice cultivation
- Sustainable razing
- Avoided grassland conversion



- Algae & Plankton
- Sediments of mangroves, salt marshes and seagrasses

Based on data from Proceedings of the National Academy of Sciences of the United States of America

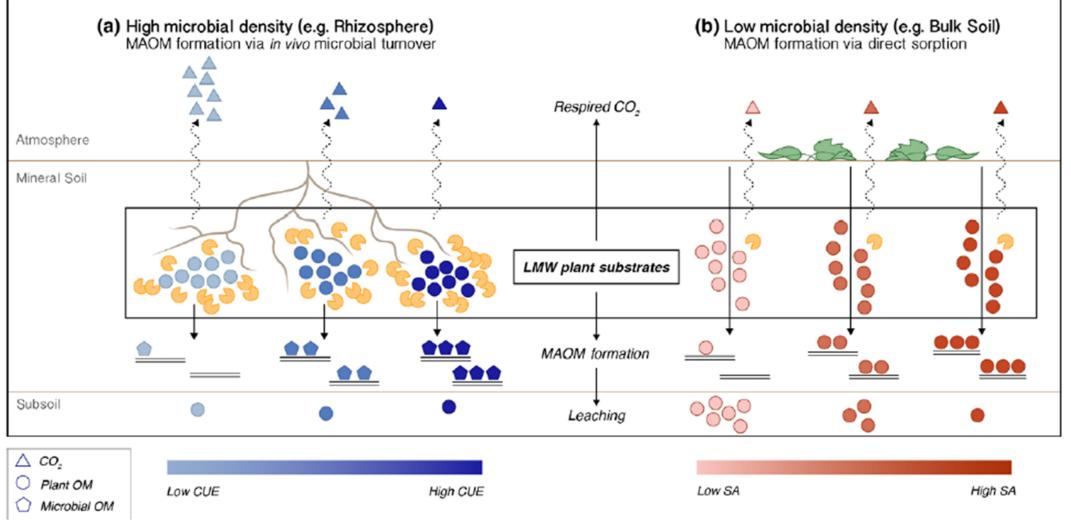




Source: Griscom et al. (PNAS 2017)



Microbes drive C storage in soil



Source: Sokol et al. (Global Change Biology 2018)



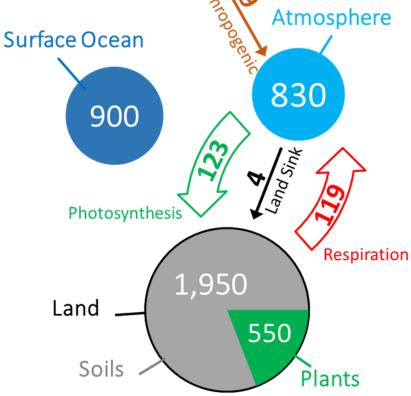
The Need for Accurate and affordable Carbon Flux Measurements

Facts of today:

- Land sink is globally significant: ~ 48 % of anthropogenic emissions.
- 2. The mean flux is small: 4 PgC yr¹ globally \approx 27 gC m⁻² yr¹
- 3. Monitoring, Reporting and Verification (MRV) must accurately resolve small fluxes.
- Current MRV methodologies are a large cost of every NBS activity

NBS challenge:

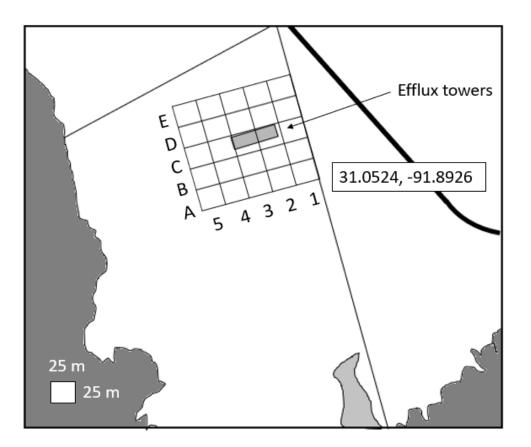
- Create affordable and auditable quantification of net CO₂ flux to support and verify NBS activities.
- 2. Reducing MRV costs to expand NBS mitigation options.



Based on IPCC AR5 Global Carbon Pools and Fluxes UNITS: Pools: PgC Fluxes: PgC yr¹



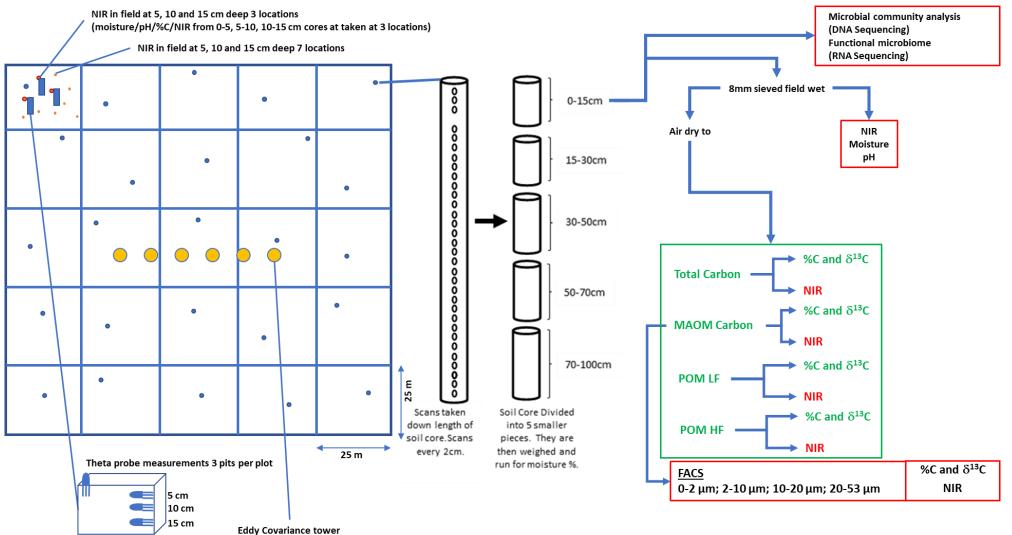
Field site







Sampling for carbon



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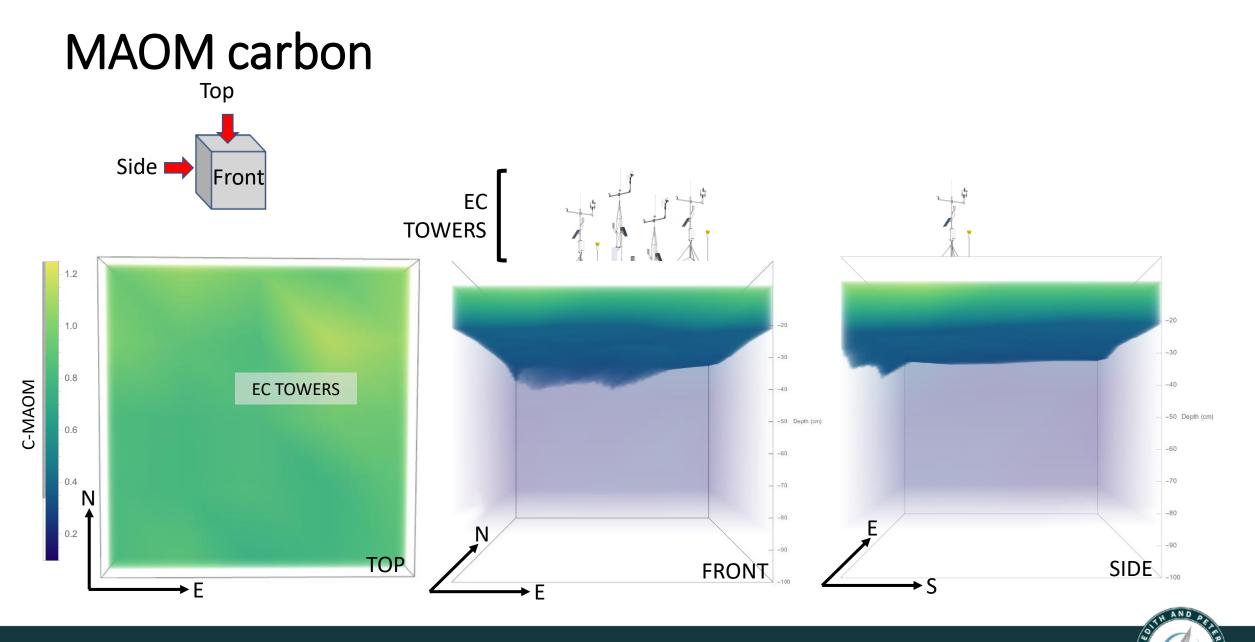
AND

NELL A

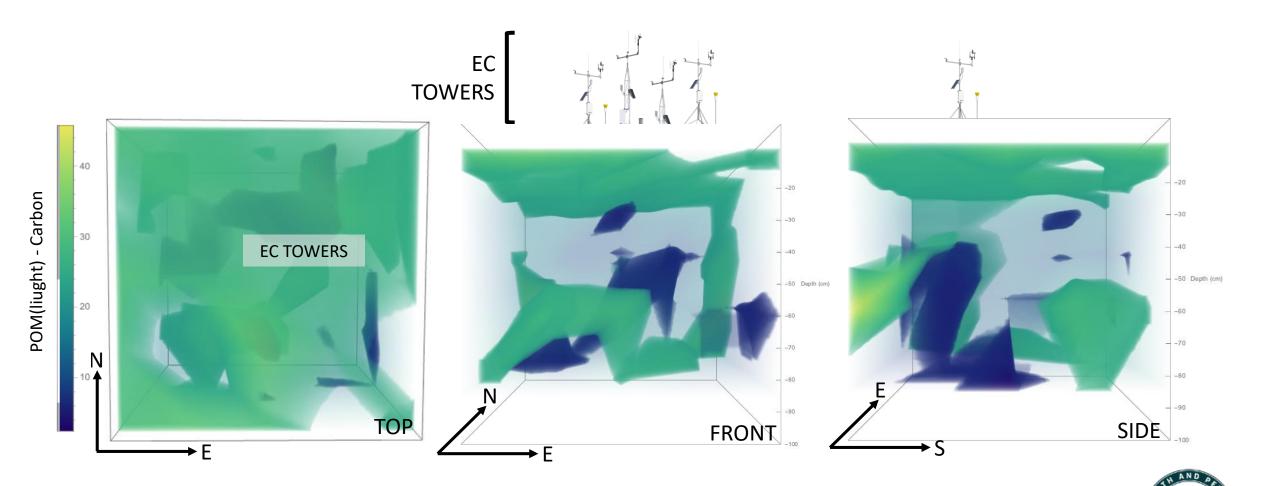
DNA Sequencing

- Purified DNA was prepared from sequencing using the Oxford Nanopore Technologies (ONT) ligation preparation kit (SQK-LSK109)
- Prepared DNA was sequenced using the ONT MinION device and flowcell (R9.4a). Resulting DNA sequence reads were basecalled using a MinIT and guppy basecaller.
- The advantage of using the ONT MinION is the ability to sequence long DNA fragments, enabling greater accuracy in taxonomically classifying the microbiome.

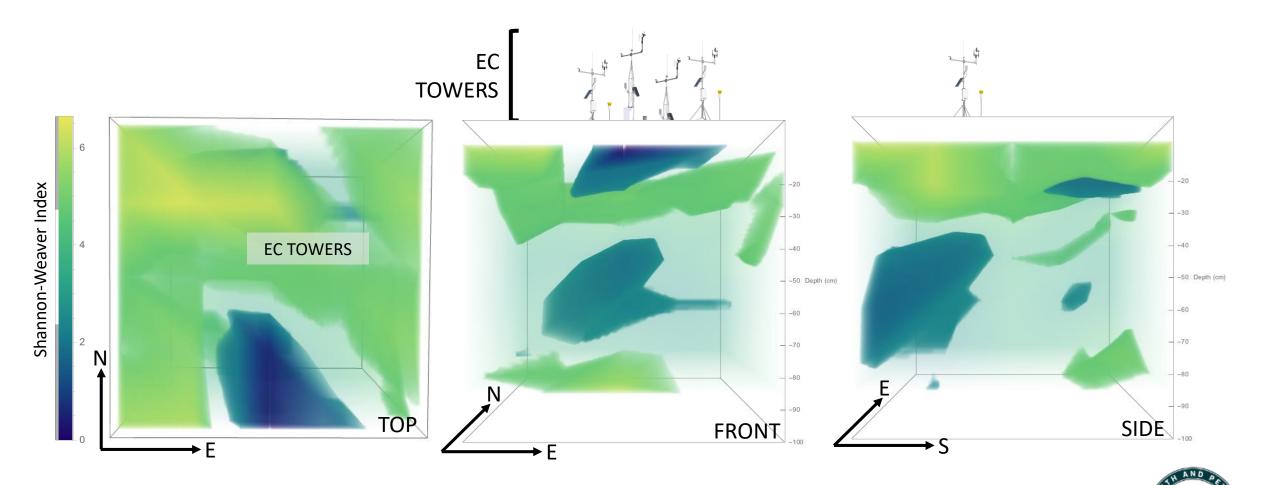




POM carbon



Alpha biodiversity



Correlations between taxa and soil chemistry

Taxa - Family	r value	Acidobacteriaceae	Bryobacteraceae	Solibacteraceae	Pyrinomonadaceae	Holophagaceae	Therm
C_MAOM	1	-0.382489632	-0.032203615	-0.348404145	0.012937625	-0.036765039	
Verrucomicrobiaceae	0.335230336	-0.116302516	0.021404272	-0.159061258	0.138050689	-0.015605714	
Gallionellaceae	0.297997208	-0.110921582	-0.021202959	-0.099689213	-0.017095328	0.017447487	
Arthrodermataceae	0.291536495	-0.196057452	-0.012247067	-0.192400181	0.045102557	-0.01449195	
llumatobacteraceae	0.271126133	-0.122234624	-0.016004617	-0.120949027	-0.004160413	-0.008847634	
Parachlamydiaceae	0.268680786	-0.138422492	0.006314389	-0.131102044	0.07739238	0.08536704	
Aerococcaceae	0.262717652	-0.098129422	0.003513034	-0.096609164	0.013685779	-0.007069418	
Gigasporaceae	0.258667176	-0.125812777	-0.023512976	-0.124113459	0.05871221	-0.009152277	
Vicinamibacteraceae	0.255873216	-0.150684262	0.012650636	-0.146909923	0.14147104	-0.011640237	
Tolypothrichaceae	0.251860196	-0.093282587	-0.01140155	-0.090519502	-0.001233811	-0.006798161	
Cytophagaceae	0.242231084	-0.084196886	-0.023396358	-0.06795507	0.002050523	-0.010535703	
Fulvivirgaceae	0.240045094	-0.103243741	-0.002548504	-0.101983717	-0.025222639	-0.007512106	
Moraxellaceae	0.232261333	-0.083700575	0.003035765	-0.082790277	-0.020511544	-0.006108991	
Syntrophaceae	0.229421123	-0.093708483	0.003771426	-0.142868329	-0.00883157	-0.013665068	
Lindgomycetaceae	0.229229355	-0.101242575	-0.009605321	-0.101213643	0.062482611	-0.007305513	
Gordoniaceae	0.228494227	-0.121306081	-0.01318444	-0.120816455	0.062868791	-0.008696237	
Teratosphaeriaceae	0.219703548	-0.134777823	0.013014443	-0.133045141	0.062626092	-0.009762387	
Cellulomonadaceae	0.219229182	-0.090309444	-0.001467719	-0.088377487	0.019297622	-0.006444107	
Rhizopodaceae	0.219058656	-0.075289244	-0.016992834	-0.072792247	-0.01824863	-0.005435023	
Clostridiaceae	0.21641673	-0.138130126	0.014196955	-0.132794928	0.00289963	-0.011120305	
Rhodobiaceae	0.215508045	-0.096648517	0.017060229	-0.095529246	0.033864218	-0.007389628	
Didymosphaeriaceae	0.2138872	-0.08037042	0.002860078	-0.079633349	-0.019324491	-0.005755449	
Mycosphaerellaceae	0.204185112	-0.102226757	0.003630357	-0.10023566	0.033478746	-0.007305513	
Rhodocyclaceae	0.203122735	-0.134122002	0.028761862	-0.124868942	0.150017356	-0.010654753	
Kineosporiaceae	0.200490573	-0.07825264	-0.005897778	-0.078305608	0.006588724	-0.005585098	
Siphoviridae	0.199578838	-0.06959223	-0.005897778	-0.070071498	0.006588724	-0.005585098	
Nitriliruptoraceae	0.199090911	-0.06450317	-0.01466435	-0.062669305	-0.015748067	-0.004690276	
Sporomusaceae	0.198612983	-0.107816212	0.009455241	-0.106916555	0.018698758	-0.008320558	
Togniniaceae	0.19794243	-0.075540282	0.022394529	-0.074220955	0.016275805	-0.005435023	
Chlorobiaceae	0.194928827	-0.089763759	-0.010807748	-0.088502966	0.019297622	-0.006444107	
Mortierellaceae	0.194156776	-0.080647334	-0.017994659	-0.079423219	0.026375319	-0.005755449	
Pseudoalteromonadaceae	0.192705773	-0.098582934	0.012051659	-0.097871	0.013685779	-0.007069418	
Leptospiraceae	0.18860556	-0.127262423	0.010278934	-0.126025045	0.015934859	0.167501864	
Sandaracinaceae	0.18855217	-0.089306057	-0.001048851	-0.086335462	0.009665016	-0.007360194	
Thermodesulfobacteriaceae	0.182647361	-0.079096613	-0.017994659	-0.079143045	0.11777494	-0.005755449	
Brucellaceae	0.17875496	-0.123194362	0.001482217	-0.121849582	0.072478769	-0.008847634	
Spirulinaceae	0.17746932	-0.057057417	-0.012687529	-0.056838639	0.050818151	-0.004058006	
Methanobacteriaceae	0.177255841	-0.090706306	-0.020147777	-0.089318579	0.060231972	-0.006444107	
Neisseriaceae	0.176737373	-0.093314339	-0.021037728	-0.091485738	0.007937887	-0.006728751	
Lythraceae	0.175959491	-0.067490919	0.020508018	-0.06761324	0.035982136	-0.004977178	
Alsobacteraceae	0.174972601	-0.064954501	-0.006166796	-0.064495876	-0.015748067	-0.004690276	



natureresearch

Check for updates

OPEN Novel cultivated endophytic Verrucomicrobiα reveal deeprooting traits of bacteria to associate with plants

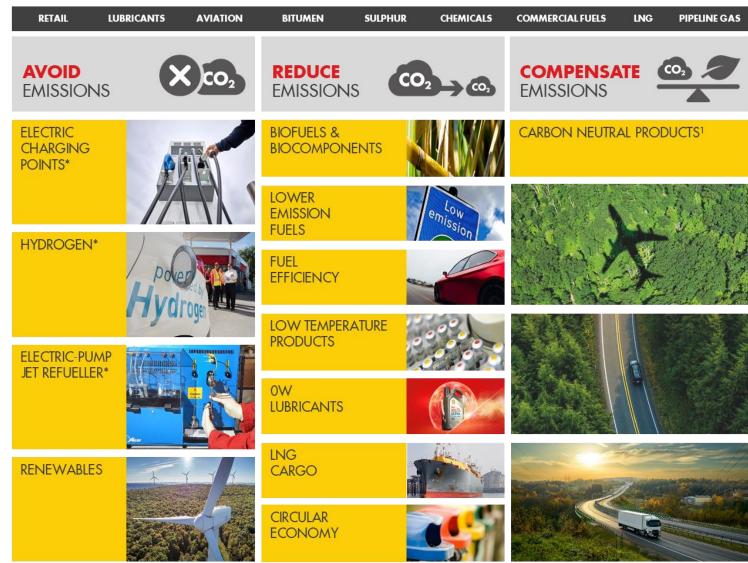
Wiebke Bünger¹, Xun Jiang¹, Jana Müller^{1,2}, Thomas Hurek¹ & Barbara Reinhold-Hurek¹

-0.005747126	-0.016485416	-0.005747126
-0.009126428	0.01643786	-0.009126428
-0.009529553	0.036234985	-0.009529553
-0.007048899	0.144357826	-0.007048899
-0.006642576	0.019718652	-0.006642576



Thriving through the energy transition, there is no single solution

To help you reduce your carbon footprint Shell offers solutions to avoid, reduce and compensate your emissions.



*When sourced from renewables

¹ The terms "Carbon Neutral", "Carbon Offset" or "Carbon off-set compensation" are applied in a non-technical way to indicate that Shell has engaged in a transaction to ensure that an amount of carbon dioxide equivalent to that associated with the production, delivery and usage of the fuel has been removed from the atmosphere through a nature-based process or emissions saved through avoided deforestation

How do nature-based solutions work?

Stabilise global warming to below 2°C



- Nature-based solutions are projects which protect or redevelop natural ecosystems, such as forests, grasslands and wetlands.
- Such nature projects can lead to the creation, marketing, trading and sale of carbon credits.
- A carbon credit represents the avoidance or removal of 1 tonne of carbon dioxide.

How much CO₂ is stored in wood?



A tree converts an average of 1 tonne of CO_2 into 1.1 cubic meters of wood. This means one cubic metre of wood contains 0.9 tonnes of CO_2 .

How fast do trees mature?

EARLY GROWTH

Trees in a forest have slow early growth, followed by a spike in growth and later growth decreases slowly. Peak growth is around 15 years for poplars and 90-100 years for oaks.

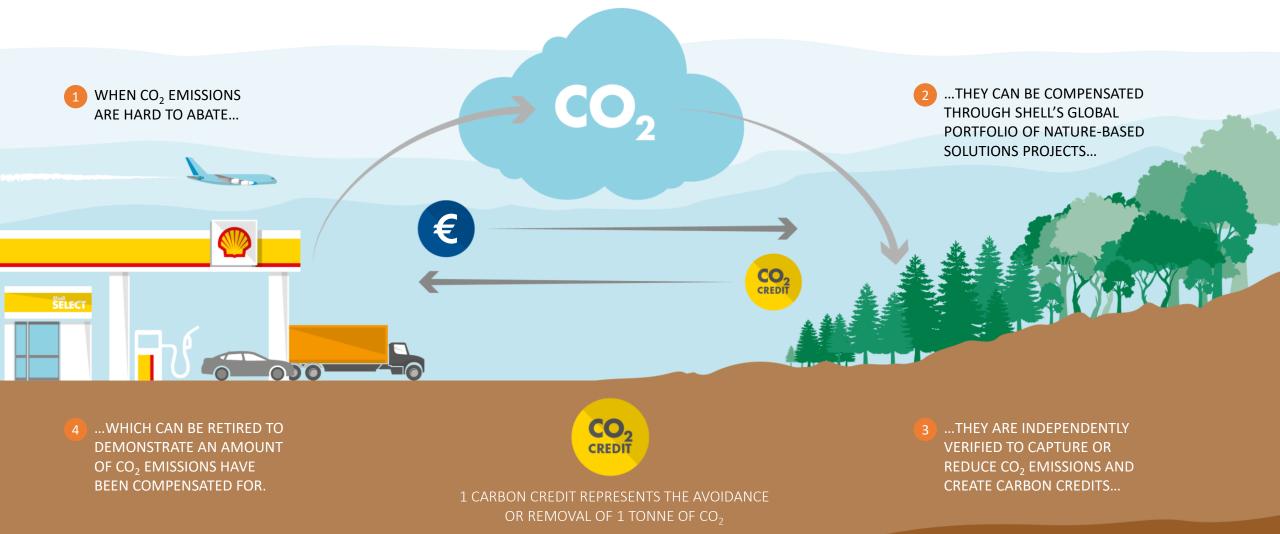
PEAK GROWTH

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LATE GROWTH



Compensate your CO₂ emissions



The fundamental criteria for the integrity of our nature-based carbon credits

PERMANENCE

If forests are cut down or destroyed, the offsets that were generated by that forest would be compromised.

NO LEAKAGE

If there would be increases in emissions caused by a project outside of its boundaries this is accounted for.

ADDITIONALITY

Emissions reductions and removals are additional to what would have happened in the absence of the project (baseline).



Our projects are assessed using approved greenhouse gas accounting methodologies

BASELINE

A baseline is established defining what would have happened in the absence of the project.

REAL

All emissions reductions and removals can be proven to have genuinely taken place.

MEASURABLE

All emissions are quantifiable, using recognised measurement tools and methodology.

UNIQUE

No more than one carbon credit can be associated with a single emission reduction of one tonne CO₂.

INDEPENDENTLY VERIFIED

All projects are verified by an independent, accredited third-party auditor.

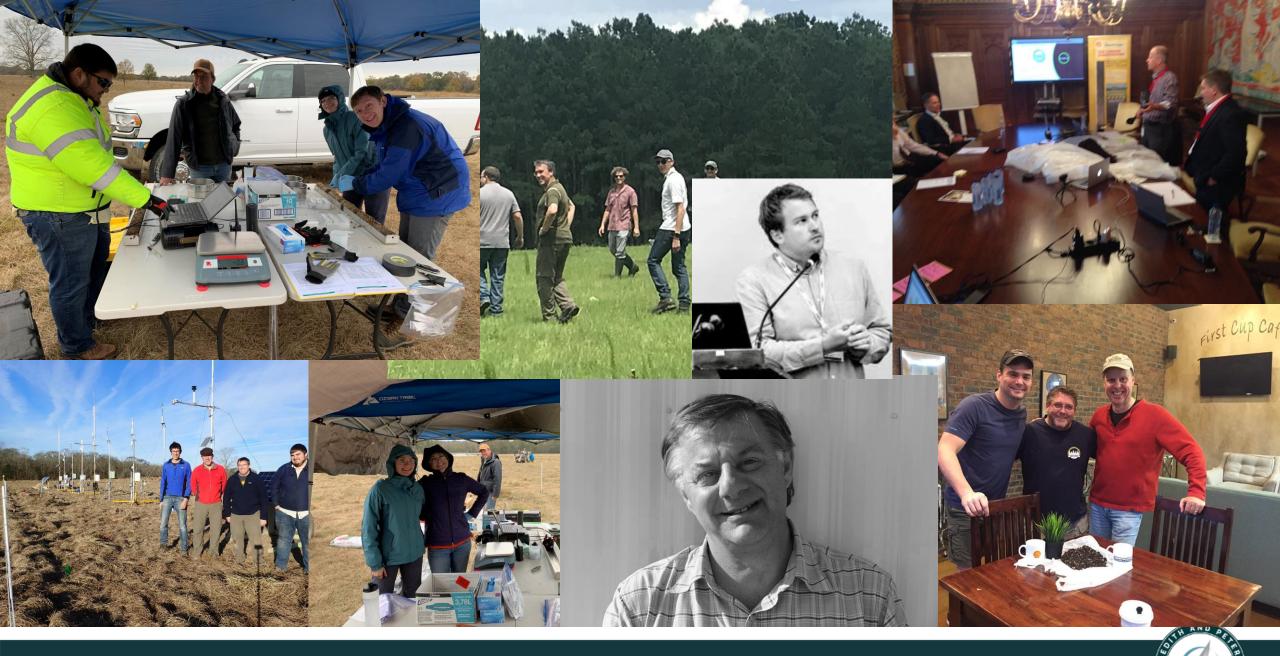
EX-POST CREDITING

Credits are only issued after the emissions reduction or removal has been verified.









NELL A