

# Isotope Tracking

## Applications

After a new product application at time  $t + 1$ , the top soil carbon isotope signature ( $\delta^{13}C$ ) for a soil layer  $k$ , is updated by balancing mass terms. Note that for pesticide applications only the first layer ( $k = z_0$ ) is considered, such that:

$$\delta^{13}C_{k(t+1)} = \frac{1}{M_{k,tot(t+1)}} \left( \delta^{13}C_{k(t)} \cdot M_{k(t)} + \delta^{13}C_{app(t+1)} \cdot M_{app(t+1)} \right)$$

$$M_{k,tot(t+1)} = M_{k(t)} + M_{app(t+1)}$$

where  $M_{k(t)}$  is the pesticide mass ( $\mu g$ ) for the layer  $k$  present before application  $app$ .

## Non-reactive transport

For each non-fractionating mass transfer process  $\delta^{13}C$  is updated also by balancing mass terms for each cell:

$$\delta^{13}C_{k(t+1)} = \frac{1}{M_{k,tot(t+1)}} \left( \delta^{13}C_{k(t)} \cdot M_{k(t)} + \delta^{13}C_{gain(t+1)} \cdot M_{gain(t+1)} - \delta^{13}C_{loss(t+1)} \cdot M_{loss(t+1)} \right)$$

$$M_{k,tot(t+1)} = M_{k(t)} + M_{gain(t+1)} - M_{loss(t+1)}$$

Update at each cell for each layer is computed by the following function, where the relevant layer and processes is selected:

```
def update_layer_delta(model, layer, process, mass_process, mass_before_transport):
    if layer == 0:
        delta_layer = model.delta_z0
        delta_layer_above = None
        mass_layer = model.pestmass_z0
    elif layer == 1:
        delta_layer = model.delta_z1
        delta_layer_above = model.delta_z0
        mass_layer = model.pestmass_z1
    elif layer == 2:
        delta_layer = model.delta_z2
        delta_layer_above = model.delta_z1
        mass_layer = model.pestmass_z2

    if process == "volat":
        pass
    elif process == "runoff":
        pass
    elif process == "leach":
        pass
    elif process == "latflux":
        pass
    else:
        raise NotImplementedError
```

```
return "updated delta for delta_layer"
```

For brevity, only two examples are shown. For volatilization, the process process=“**volat**” is chosen and the updated  $\delta^{13}C$  for a soil layer  $k$  is returned as,

```
def update_layer_delta(model, layer, process, mass_process, mass_before_transport):
    if layer == 0:
        delta_layer = model.delta_z0
        delta_layer_above = 0
        mass_layer = model.pestmass_z0
    elif layer == 1:
        delta_layer = model.delta_z1
        delta_layer_above = model.delta_z0
        mass_layer = model.pestmass_z1
    elif layer == 2:
        delta_layer = model.delta_z2
        delta_layer_above = model.delta_z1
        mass_layer = model.pestmass_z2

    if process == "volat":
        mass_loss = mass_process["mass_loss"]
        mass_gain = 0
        delta_gain = 0
        delta_loss = delta_layer

    elif process == "runoff":
        pass
    elif process == "leach":
        pass
    elif process == "latflux":
        pass
    else:
        raise NotImplementedError

    if process == "latflux":
        pass
    else:
        mass_tot = mass_before_transport + mass_gain - mass_loss
        delta_int = ((1/mass_tot) *
                     (delta_layer * mass_before_transport + # initial
                      delta_gain * mass_gain - # mass_in
                      delta_loss * mass_loss)) # mass_out

    return delta_int
```

For lateral flux, update follows the same approach as that for water mass exchange across cells. The pesticide mass at cell  $j$  after lateral flux is given by:

$$M_{j,tot(t+1)} = M_{j(t)} + \sum_{i=1}^{N(t)} M_{loss,i(t)} - M_{loss,j(t)}$$

The mass gain at cell  $j$  is given by the mass from upstream cells  $i$  contributing to downstream cells and given by:

$$\sum_{i=1}^{N(t)} M_{loss,i(t)} = \frac{W_j \sum_{i=1}^{N(t)} \max[C_{i,aq} \cdot (c_z(SW_i - SW_{fc,i})), 0]}{\sum_{i=1}^{N(t)} W_i}$$

Loss at cell  $j$  is then given by:

$$M_{loss,j(t)} = C_{j,aq} \cdot (c_z(SW_j - SW_{fc,j}))$$

Considering the isotope mass balance, while simplfying for the relative wetness index at cell  $j$ , we obtain:

$$W_{j/i} = \frac{W_j}{\sum_{i=1}^{N(t)} W_i}$$

$$\delta^{13}C_{j(t+1)} = \frac{1}{M_{j,tot(t+1)}} \left( \delta^{13}C_{j(t)} \cdot M_{j(t)} + W_{j/i} \sum_{i=1}^{N(t)} \max[\delta^{13}C_i \cdot C_{i,aq} \cdot (c_z(SW_i - SW_{fc,i})), 0] - \delta^{13}C_{j(t)} \cdot M_{loss,j(t)} \right)$$

The `update_layer_delta` functions, takes process=“**latflux**” and implements the above equations as:

```
def update_layer_delta(model, layer, process, mass_process, mass_before_transport):
    if layer == 0:
        delta_layer = model.delta_z0
        delta_layer_above = 0
        mass_layer = model.pestmass_z0
    elif layer == 1:
        delta_layer = model.delta_z1
        delta_layer_above = model.delta_z0
        mass_layer = model.pestmass_z1
    elif layer == 2:
        delta_layer = model.delta_z2
        delta_layer_above = model.delta_z1
        mass_layer = model.pestmass_z2

    if process == "volat":
        pass
    elif process == "runoff":
        pass
    elif process == "leach":
        pass
    elif process == "latflux":
        mass_latflux = mass_process["net_mass_latflux"] # mg
        mass_loss = mass_process["cell_mass_loss_downstream"]
        mass_gain = mass_process["upstream_mass_inflow"]
        mass_tot = mass_before_transport + mass_gain - mass_loss
        # Proof of first fraction, i.e., f1 > 1
        f1 = mass_before_transport / mass_tot
        # Proof of 2nd (inflow) fraction, f2 < 1
        # f2 = accuflux(model.ldd, mass_gain)/accuflux(model.ldd, mass_tot)
        f3 = mass_loss / mass_tot # Proof of third (leaving) mass fraction, f3 < 1
    else:
        raise NotImplementedError
    if process == "latflux":
        delta2_f2 = accuflux(model.ldd_subs, mass_gain*delta_layer)/accuflux(model.ldd_subs, mass_tot)
```

```

        delta_int = (delta_layer * f1) + delta2_f2 - (delta_layer * f3)
    else:
        pass

    return delta_int

```

Full function

This is my change.

```

def update_layer_delta(model, layer, process, mass_process, mass_before_transport):
    if layer == 0:
        delta_layer = model.delta_z0
        delta_layer_above = 0
        mass_layer = model.pestmass_z0
    elif layer == 1:
        delta_layer = model.delta_z1
        delta_layer_above = model.delta_z0
        mass_layer = model.pestmass_z1
    elif layer == 2:
        delta_layer = model.delta_z2
        delta_layer_above = model.delta_z1
        mass_layer = model.pestmass_z2
    if process == "volat":
        mass_loss = mass_process["mass_loss"]
        mass_gain = 0
        delta_gain = 0
        delta_loss = delta_layer
    elif process == "runoff":
        mass_loss = mass_process["mass_runoff"]
        mass_gain = 0
        delta_gain = 0
        delta_loss = delta_layer
    elif process == "leach":
        mass_leached = mass_process["mass_leached"] # mg
        mass_loss = mass_leached
        mass_gain = 0
        delta_gain = delta_layer_above
        delta_loss = delta_layer
    elif process == "latflux":
        mass_latflux = mass_process["net_mass_latflux"] # mg
        mass_loss = mass_process["cell_mass_loss_downstream"]
        mass_gain = mass_process["upstream_mass_inflow"]
        mass_tot = mass_before_transport + mass_gain - mass_loss
        # Proof of first fraction, i.e., f1 > 1
        f1 = mass_before_transport / mass_tot
        # Proof of 2nd (inflow) fraction, f2 < 1
        # f2 = accuflux(
        #     model.ldd, mass_gain)/accuflux(model.ldd, mass_tot)
        # Proof of third (leaving) mass fraction, f3 < 1
        f3 = mass_loss / mass_tot
    else:
        raise NotImplementedError
    if process == "latflux":

```

```

    delta2_f2 = accuflux(model.1dd_subs, mass_gain*delta_layer)/accuflux(model.1dd_subs, mass_tot)
    delta_int = (delta_layer * f1) + delta2_f2 - (delta_layer * f3)
else:
    mass_tot = mass_before_transport + mass_gain - mass_loss
    delta_int = ((1/mass_tot) *
                  (delta_layer * mass_before_transport + # initial
                   delta_gain * mass_gain - # mass_in
                   delta_loss * mass_loss)) # mass_out
# return {"delta_int": delta_int, "mass_layer": mass_layer}
return delta_int

```

## References