## **Equations S3**. GMA mass balance equations for labelad pools.

$$\begin{split} dL_1/dt &= l_{1,2,1} - l_{1,2} \\ dL_2/dt &= \left(l_{1,2} + l_{3,2} + l_{4,2}\right) - \left(l_{2,3}^a + l_{2,4} + l_{2,5}\right)^a \\ dL_3/dt &= \left(l_{2,3}^c + l_{8,3} + l_{18,3} + l_{19,3}\right) - \left(l_{3,2} + l_{3,7} + l_{3,8}^a\right) \\ dL_4/dt &= l_{2,4} - \left(l_{4,2} + l_{4,17}\right) \\ dL_5/dt &= \left(l_{2,5} + l_{6,5} + l_{7,5}\right) - \left(l_{5,6} + l_{5,7}^a\right) \\ dL_6/dt &= l_{5,6} - \left(l_{6,5} + l_{6,17}\right) \\ dL_7/dt &= \left(l_{3,7} + l_{5,7}^c + l_{8,7} + l_{18,7} + l_{19,7}\right) - \left(l_{7,5} + l_{7,8}^a + l_{7,43}\right) \\ dL_8/dt &= \left(l_{6,8}^c + l_{6,8}^c + l_{20,8}\right) - \left(l_{8,3} + l_{8,7}^c + l_{8,18} + l_{8,20}\right) \\ dL_9/dt &= l_{11,9} - \left(l_{9,10} + l_{9,15}\right) \\ dL_{10}/dt &= l_{1,19} - \left(l_{11,9} + l_{11,14}\right) \\ dL_{12}/dt &= \left(l_{4,17} + l_{6,17} + l_{24,12}^c\right) - \left(l_{12,1} + l_{12,11} + l_{12,23}^a + l_{12,148}\right) \\ dL_{14}/dt &= \left(l_{5,8}^b + l_{7,8}^b + l_{11,14} + l_{18,19}^b\right) - \left(l_{14,12} + l_{14,145}\right) \\ dL_{15}/dt &= l_{9,15} - \left(l_{5,8}^b + l_{7,8}^b + l_{15,44} + l_{18,19}^b\right) \\ dL_{17}/dt &= \left(l_{4,17} + l_{6,17}\right) - l_{14,145} \\ dL_{18}/dt &= \left(l_{8,18} + l_{21,18}\right) - \left(l_{18,3} + l_{18,7} + l_{18,19}^a + l_{18,21}\right) \\ dL_{29}/dt &= l_{8,20} - l_{20,8} \\ dL_{21}/dt &= l_{18,21} - l_{21,18} \\ dL_{22}/dt &= l_{19,22} - l_{22,19} \\ dL_{23}/dt &= \left(l_{2,23} - \left(l_{12,3}^b + l_{5,7}^b\right) \\ dL_{24}/dt &= l_{25,24} - \left(l_{12,23}^b + l_{24,12}^b\right) \\ dL_{26}/dt &= l_{25,26} - l_{26,27} \\ dL_{27}/dt &= l_{26,27} - l_{27,28} \\ dL_{29}/dt &= l_{28,29} - l_{29,30} \\ dL_{30}/dt &= \left(l_{20,30} + l_{33,30}\right) - \left(l_{30,31} + l_{30,33}\right) \\ dL_{30}/dt &= \left(l_{31,32} + l_{35,32} + l_{37,32} + l_{39,32}\right) - \left(l_{32,35} + l_{32,37} + l_{32,39} + l_{32,186}\right) \\ dL_{35}/dt &= \left(l_{31,32} + l_{35,32} + l_{37,32} + l_{39,32}\right) - \left(l_{32,35} + l_{32,37} + l_{32,39} + l_{32,186}\right) \\ dL_{36}/dt &= \left(l_{31,34} - l_{34,31}\right) - \left(l_{36,37} + l_{36,37} + l_{36,37} + l_{36,37} + l_{36,37} + l_{36,37} + l_{36,37}\right) \\ dL_{36}/dt &= \left(l_{37,36} + l_{39,36}\right) - \left(l_{36,37} + l_{36,37} + l_{36,37} + l_{36,37} + l_{36,37} + l_{36,37}\right) \\ dL_{36}/dt &= \left(l_{37,36} + l_{39,36}\right$$

$$\begin{split} dL_{37}/dt &= \left(l_{32,37} + l_{36,37}^a + l_{36,37}^b + l_{36,37}^c\right) - \left(l_{37,32} + l_{37,36}\right) \\ dL_{38}/dt &= l_{125,38} - l_{38,25} \\ dL_{39}/dt &= \left(l_{42,39} + l_{36,39} + l_{40,39}\right) - \left(l_{39,32} + l_{39,36}\right) \\ dL_{40}/dt &= l_{35,40} - \left(l_{40,35} + l_{40,39}\right) \end{split}$$

(\*) A superscript indicates deviations from the total pool; the method automatically computes equations for the labeled fractions.

## Tracer method: Mathematical approach to modeling tracer dynamics

To characterize the movement and distribution of a radioactive tracer throughout the sphingolipid pathway, simulations were performed using a method specifically developed for this purpose [1]. This method models the dynamics of total (labeled plus unlabeled) metabolites and in a second and third set of equations distributes material between labeled and unlabeled fractions (see [1] and main manuscript for details of the method).

## **Tracer equations**

In Equations S3 and S4, the labeled metabolites serine ( $L_{13}$ ) and inositol ( $L_{16}$ ) are not included because they never receive label in radioactive acetate ( $L_{125}$ ) experiments. For the same reason, the time independent variables palmitate ( $L_{158}$ ) and pyruvate ( $L_{124}$ ) are not included in Equations S3 and S4.

## References.

1. Voit EO, Alvarez-Vasquez F, Sims KJ (2004) Analysis of dynamic labeling data. Math Biosci 191: 83-99.