## Accurate and Efficient Solution of Linear and Nonlinear ME Models

## The need for quadruple precision

Carrying somewhat more precision in the arithmetic than twice the precision carried in the data and available for the result will vastly reduce embarrassment due to roundoff-induced anomalies
Default evaluation in Quad is the humane option

- William Kahan (2012)


## Quad datatypes are now available

GCC provides real (16) and float128 in gfortran and C, C++ This is the humane option for producing quad software

We use double-MINOS and quad-MINOS (f77 sparse LP/NLP) called from f77, f90, or python

## Metabolic networks

flux balance analysis (FBA)
Constraint-based models enable the study of metabolism at genome-scale

- M models: multiscale reconstructions of metabolism
- ME models include protein expression (even more multiscale)
- Stoichiometric matrix $S$, fluxes $v$, growth rate $\mu$

Most coefficients are moderate: $S_{i j}=0, \pm 1, \pm 2$
Some coefficients are large: $\quad S_{i j}=10,000$

- Similarly for fluxes because of coupling constraints:

$$
v_{i} / v_{j} \geq \mu / k_{\text {eff }} \quad \Rightarrow \quad v_{i} \geq\left(\mu / k_{\text {eff }}\right) v_{j}
$$

Models are linear when $\boldsymbol{\mu}$ is fixed

$$
\begin{aligned}
& \max \boldsymbol{c}^{\boldsymbol{T}} \boldsymbol{v} \\
& \text { st } \boldsymbol{S} \boldsymbol{v}=\boldsymbol{b} \\
& \quad \text { bounds on } \boldsymbol{v}
\end{aligned}
$$

Solved by

- openCOBRA toolbox (CPLEX, glpk, Gurobi, Mosek, MINOS, ...)
- MONGOOSE toolbox (exact simplex solver QSOpt_ex)

- Exact simplex as in MONGOOSE can handle nonlinear $\boldsymbol{\mu}$ via binary search, but is not scalable to ME models
- Quad-precision LP and NLP provide a balance between reliability and speed

Linear ME model of E. coli double-MINOS, quad-MINOS LP
Problem GlcAerWT (Thiele, Fleming, et al. 2012), $68300 \times 76664$ Step 1: double-MINOS, cold start, scaling


Step 2: quad-MINOS, warm start, scaling


Step 3: quad-MINOS, warm start, no scaling


tinyME (Yang et al. 2014), $2512 \times 2828$


Nonlinear ME model
solveME (Yang et al. 2015), $11386 \times 18755$


## References

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