

The DQQ procedure for multiscale optimization

Ding Ma and Michael Saunders

MS&E and ICME, Stanford University

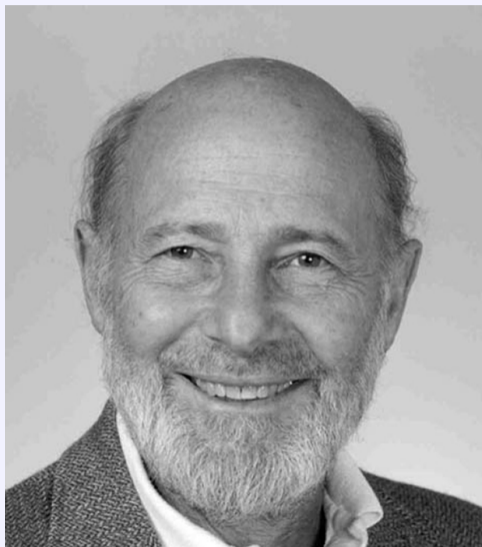
ICCOPT 2016

Tokyo, Japan, August 6–11, 2016

Scientific Computing and Matrix Computations Seminar

UC Berkeley, Wed Oct 12, 2016

Prof Joseph Keller, 1923–2016



SCMC Seminar, UC Berkeley, Oct 12, 2016

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Inverse problems

Q: ?

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Nonmonotonic world records

Google search: ~~Missing: nonmonotonic~~

William Kahan, LA/Opt seminar, Oct 13, 2011

Desperately Needed Remedies for the Undebuggability of Large Floating-Point Computations in Science and Engineering



SCMC Seminar, UC Berkeley, Oct 12, 2016

Coauthor Ding Ma at INFORMS SFO, Dec 2014



ICCOPT Tokyo, Aug 2016



ICCOPT Tokyo, Aug 2016



ICCOPT Tokyo, Aug 2016



Bus tour to Mt Fuji



Mt Fuji



Mt Fuji



Abstract

Constrained optimization solvers typically scale the constraints, solve the scaled problem, then unscale. With multiscale problems and a conventional double-precision solver, **the unscaled solution may not satisfy the constraints well**. Our DQQ procedure continues with a quad-precision version of the solver to obtain accurate solutions efficiently.

A prime application is to **metabolic networks in systems biology**. Keywords are flux balance analysis (FBA) and genome-scale modeling of Metabolism and macromolecular Expression (ME models). DQQ typically achieves at least 20 digits of precision. (Joint work with Ding Ma)

For smooth functions with few variables, Quad improves the performance of **quasi-Newton optimization with finite-difference gradients**. We illustrate with IMSPE-optimal design of computer experiments. (Joint work with Selden Cray)

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- 6 Multiscale NLPs
- 7 Finite-difference gradients
- 8 Conclusions

Motivation

Multiscale LPs in systems biology

$$\max_v c^T v \quad \text{s.t.} \quad Sv = 0, \quad l \leq v \leq u$$

Normal for LP (simplex or interior)

- Scale to reduce large S_{ij}
- Solve with Feasibility/Optimality tols $1e-6$
- Unscale

Multiscale LPs in systems biology

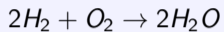
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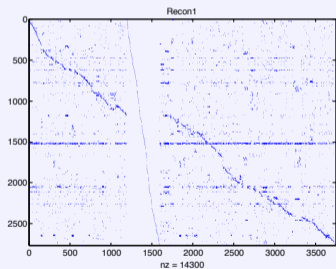
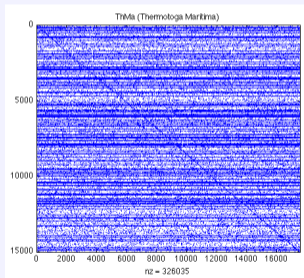
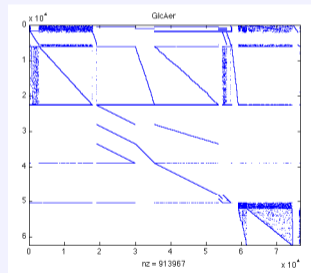
- Scale to reduce large S_{ij}
- Solve with Feasibility/Optimality tols $1e-6$
- Unscale

Difficulty

- Unscaling magnifies residuals
- Solution may be far from feasible or optimal

Stoichiometric matrices S 

	·	·	·	W	·	·	·
H_2	·	·	·	-2	·	·	·
O_2	·	·	·	-1	·	·	·
H_2O	·	·	·	2	·	·	·

chemicals \times reactions2800 \times 370015000 \times 1800062000 \times 77000

ME models (Metabolism and macromolecular Expression)

Coupling constraints

$$c_{\min} \leq \frac{v_i}{v_j} \leq c_{\max} \quad \equiv \quad c_{\min} v_j \leq v_i, \quad v_i \leq c_{\max} v_j$$

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Lifting

$$v_1 \leq 10000 v_2 \quad \equiv \quad v_1 \leq 100 s_1, \quad s_1 \leq 100 v_2$$

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Iterative refinement for LP problems

Ambros M. Gleixner (2015).

Exact and Fast Algorithms for Mixed-Integer Nonlinear Programming,
PhD thesis, ZIB, Technical University of Berlin.

Compute residuals in multiple precision, solve LP problem for correction, repeat.

Quad 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*

Methods for achieving Quad precision

Double auxiliary functions (hand-coded)

Even `qdotdd(v,w)` needs `twosum`, `split`, `twoproduct` `sum2`, `dot2`
to compute \mathbf{x}, \mathbf{y} such that $\mathbf{q} = \mathbf{v}^T \mathbf{w} = \text{quad}(\mathbf{x}) + \text{quad}(\mathbf{y})$

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Double-double datatype (≈ 32 digits)

QD: <http://crd-legacy.lbl.gov/~dhbailey/mpdist/>

Minor changes to source code (except I/O), uses floating-point hardware

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Minor changes to source code (except I/O), uses floating-point hardware

Quad datatype (≈ 34 digits)

`real(16)` in gfortran, ifort `float128` in gcc

Minor changes

We use this humane approach to Quad implementation

Double MINOS and Quad MINOS

real(8)

eps = 2.22e-16

Hardware

real(16)

eps = 1.93e-34

Software

snopt9 = Double or Quad SQOPT, SNOPT

snPrecision.f90

```
module snModulePrecision
  implicit none
  public
  integer(4), parameter :: ip = 4, rp = 8    ! double
  ! integer(4), parameter :: ip = 8, rp = 16 ! quad
end module snModulePrecision
```

snopt9 = Double or Quad SQOPT, SNOPT

snPrecision.f90

```
module snModulePrecision
  implicit none
  public
  integer(4), parameter :: ip = 4, rp = 8    ! double
  ! integer(4), parameter :: ip = 8, rp = 16 ! quad
end module snModulePrecision
```

module sn50lp

```
use snModulePrecision, only : ip, rp
subroutine s5solveLP ( x, y )
real(rp), intent(inout) :: x(nb), y(nb)
```

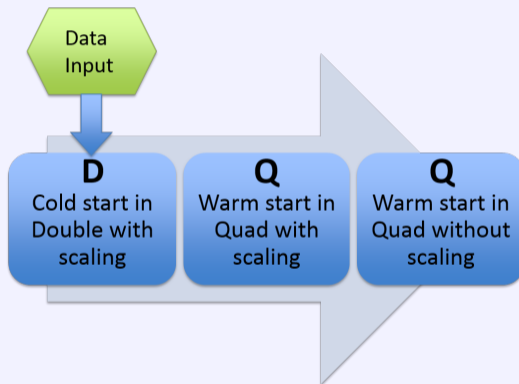
DQQ procedure

Step D: Double MINOS

Step Q1: Quad MINOS

Step Q2: Quad MINOS with no scaling

DQQ procedure



MINOS runtime options for DQQ procedure

	Default	Step D	Step Q1	Step Q2
Scale option	2	2	2	0
Feasibility tol	1e-6	1e-7	1e-15	1e-15
Optimality tol	1e-6	1e-7	1e-15	1e-15
Expand frequency	10000	100000	100000	100000
LU Factor tol	100.0	1.9	10.0	5.0
LU Update tol	10.0	1.9	10.0	5.0

pilot models from Netlib
 Mészáros *problematic* LPs
 ME biochemical network models

	m	n	$\text{nnz}(A)$	$\max A_{ij} $	$\ x^*\ _\infty$	$\ y^*\ _\infty$
pilot4	411	1000	5145	3e+04	1e+05	3e+02
pilot	1442	3652	43220	2e+02	4e+03	2e+02
pilot87	2031	4883	73804	1e+03	2e+04	1e+01
de063155	853	1488	5405	8e+11	3e+13	6e+04
de063157	937	1488	5551	2e+18	2e+17	6e+04
de080285	937	1488	5471	1e+03	1e+02	3e+01
gen1	770	2560	64621	1e+00	3e+00	1e+00
gen2	1122	3264	84095	1e+00	3e+00	1e+00
gen4	1538	4297	110174	1e+00	3e+00	1e+00
l30	2702	15380	64790	1e+00	1e+09	4e+00
iprob	3002	3001	12000	1e+04	3e+02	1e+00
TMA_ME	18210	17535	336302	2e+04	6e+00	1e+00
GlcAerWT	68300	76664	926357	8e+05	6e+07	2e+07
GlcAlift	69529	77893	928815	3e+05	6e+07	2e+07

Itns and runtimes in secs for DQQ Steps 1,2,3 (Double MINOS and Quad MINOS).

Pinf and Dinf = \log_{10} max primal/dual infeasibilities.

	Itns	Times	Final objective	Pinf	Dinf
pilot4	1571	0.1	-2.5811392602e+03	-05	-13
	6	0.0	-2.5811392589e+03	-39	-31
	0	0.0	-2.5811392589e+03	-	-30
pilot	16060	5.7	-5.5739887685e+02	-06	-03
	29	0.7	-5.5748972928e+02	-	-27
	0	0.2	-5.5748972928e+02	-	-32
pilot87	19340	15.1	3.0171038489e+02	-09	-06
	32	2.2	3.0171034733e+02	-	-33
	0	1.2	3.0171034733e+02	-	-33

Step 3 Pinf/ $\|x^*\|_\infty$ and Dinf/ $\|y^*\|_\infty$ are $\mathbf{O}(10^{-30})$

	Itns	Times	Final objective	Pinf	Dinf
de063155	921	0.0	1.8968704286e+10	-13	+03
	78	0.1	9.8830944565e+09	-	-17
	0	0.0	9.8830944565e+09	-	-24
de063157	488	0.0	1.4561118445e+11	+20	+18
	476	0.5	2.1528501109e+07	-27	-12
	0	0.0	2.1528501109e+07	-	-12
de080285	418	0.0	1.4495817688e+01	-09	-02
	132	0.1	1.3924732864e+01	-35	-32
	0	0.0	1.3924732864e+01	-	-32
gen1	369502	205.3	-1.6903658594e-08	-06	-12
	246428	9331.3	1.2935699163e-06	-12	-31
	2394	81.6	1.2953925804e-06	-45	-30
gen2	44073	60.0	3.2927907828e+00	-04	-11
	1599	359.9	3.2927907840e+00	-	-29
	0	10.4	3.2927907840e+00	-	-32
gen4	45369	212.4	1.5793970394e-07	-06	-10
	53849	14812.5	2.8932268196e-06	-12	-30
	37	10.4	2.8933064888e-06	-54	-30

	Itns	Times	Final objective	Pinf	Dinf
l30	1229326	876.7	9.5266141574e-01	-10	-09
	275287	7507.1	-7.5190273434e-26	-25	-32
	0	0.2	-4.2586876849e-24	-24	-33
iprob	1087	0.2	2.6891551285e+03	+02	-11
	0	0.0	2.6891551285e+03	+02	-31
	0	0.0	2.6891551285e+03	+02	-28
TMA_ME	12225	37.1	8.0051076669e-07	-06	-05
	685	61.5	8.7036315385e-07	-24	-30
	0	6.7	8.7036315385e-07	-	-31
GlcAerWT	62856	9707.3	-2.4489880182e+04	+04	-05
	5580	3995.6	-7.0382449681e+05	-07	-26
	4	60.1	-7.0382449681e+05	-19	-21
GlcAlift	134693	14552.8	-5.1613878666e+05	-03	-01
	3258	1067.1	-7.0434008750e+05	-09	-26
	2	48.1	-7.0434008750e+05	-20	-22

Step 3 Pinf/ $\|x^*\|_\infty$ and Dinf/ $\|y^*\|_\infty$ are $\mathbf{O}(10^{-30})$
 except **iprob** is infeasible

DRR procedure

Step D: Double MINOS

Step R1: Refinement

Step R2: Refinement with no scaling

Plausible alternative to DQQ

DRR procedure

- We need **Quad residuals** for solving $Bx_B = b - Nx_n$ after LU and for solving $Bp = a$, $B^T y = c_B$ each iteration

DRR procedure

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- **Quad** $r = a - Bp$ needs $r \leftarrow r - p_k B_k$ (qaxpy)
Compiler converts B to **Quad** every iteration

DRR procedure

- We need **Quad residuals** for solving $Bx_B = b - Nx_n$ after LU and for solving $Bp = a$, $B^T y = c_B$ each iteration
- **Quad** $r = a - Bp$ needs $r \leftarrow r - p_k B_k$ (**qaxpy**)
Compiler converts B to **Quad** every iteration
- **Quad** $r = c_B - B^T y$ needs **Quad dotproducts** (**qdot**)
Again, compiler converts B to **Quad** every iteration

LPnetlib test problems

Compare DRR and DQQ

62 classic LP problems (ordered by file size)

afiro	scfxm1	ship04s	pilotja
stocfor1	bandm	seba	ship081
adlittle	e226	grow15	nesm
scagr7	grow7	fffff800	ship121
sc205	etamacro	scfxm3	cycle
share2b	agg	ship041	greenbea
recipe	scsd1	ganges	greenbeb
vtpbase	standata	sctap2	80bau3b
share1b	beaconfd	grow22	d2q06c
bore3d	gfrdpnc	ship08s	woodw
scorpion	stair	stocfor2	d6cube
capri	scrs8	pilotwe	pilot
brandy	shell	ship12s	wood1p
scagr25	scfxm2	25fv47	pilot87
sctap1	pilot4	sierra	
israel	scsd6	czprob	

DRR procedure on LPnetlib problems

$P_{\text{inf}} = \max$ Primal infeasibility

$D_{\text{inf}} = \max$ Dual infeasibility / $(1 + \|y^*\|_{\infty})$

MINOS stops when

$P_{\text{inf}} \leq$ Feasibility tol	Default $1e-6$
$D_{\text{inf}} \leq$ Optimality tol	Default $1e-6$

Plot $\log_{10}(P_{\text{inf}})$ and $\log_{10}(D_{\text{inf}})$ for steps D, R1, R2

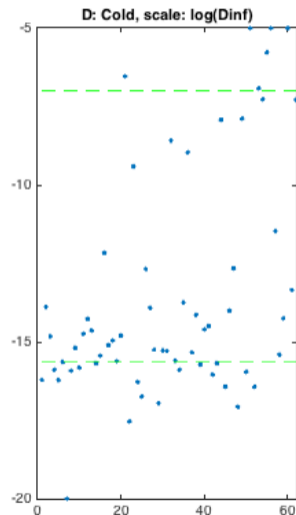
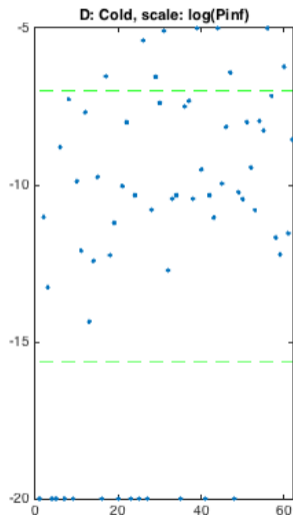
Primal/dual infeasibilities:

Step D: Double MINOS, cold start, scale

Scale option 2
 Feasibility tol $1e-7$
 Optimality tol $1e-7$

LU Partial Pivoting
 LU Factor tol 10.0
 LU Update tol 10.0
 Quad refinement 0

$\epsilon = 2.2e-16$



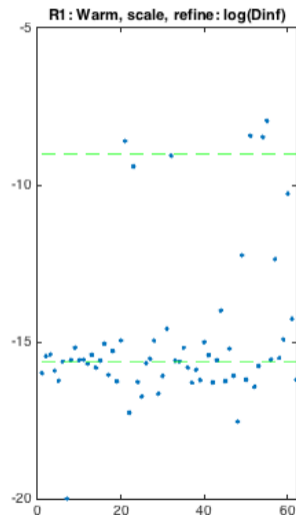
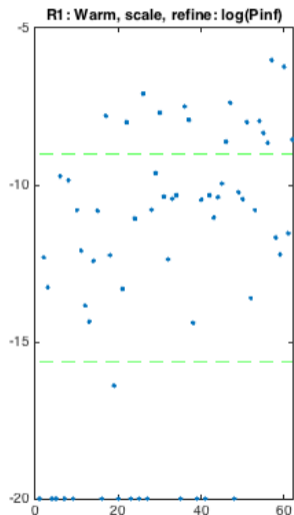
Primal/dual infeasibilities:

Step R1: Double MINOS, warm start, scale, refine

Scale option 2
 Feasibility tol 1e-9
 Optimality tol 1e-9

LU Partial Pivoting
 LU Factor tol 1.9
 LU Update tol 1.9
 Quad refinement 1

$\epsilon = 2.2e-16$



Primal/dual infeasibilities:

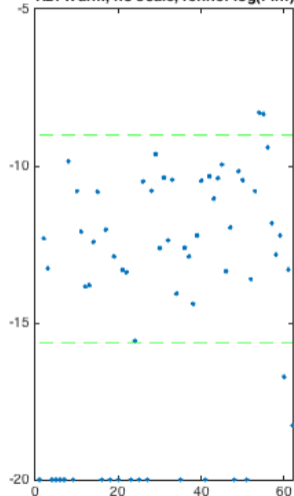
Step R2: Double MINOS, warm start, no scale, refine

```
Scale option 0
Feasibility tol 1e-9
Optimality tol 1e-9
```

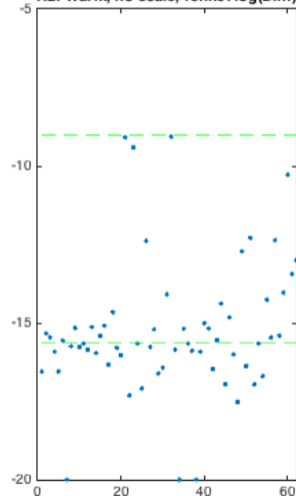
```
LU Partial Pivoting
LU Factor tol 1.9
LU Update tol 1.9
Quad refinement 1
```

 $\epsilon = 2.2e-16$

R2: Warm, no scale, refine: log(Pinf)



R2: Warm, no scale, refine: log(Dinf)



DQQ procedure on LPnetlib problems

$P_{\text{inf}} = \max$ Primal infeasibility

$D_{\text{inf}} = \max$ Dual infeasibility / $(1 + \|y^*\|_{\infty})$

MINOS stops when

$$\begin{array}{l} P_{\text{inf}} \leq \text{Feasibility tol} \\ D_{\text{inf}} \leq \text{Optimality tol} \end{array}$$

Plot $\log_{10}(P_{\text{inf}})$ and $\log_{10}(D_{\text{inf}})$ for steps D, Q1, Q2

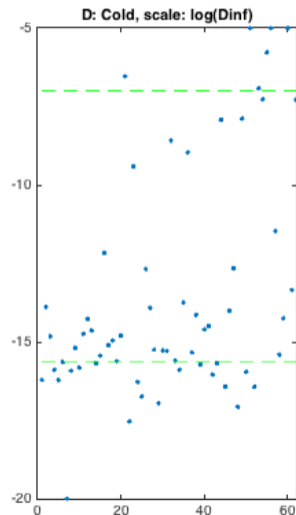
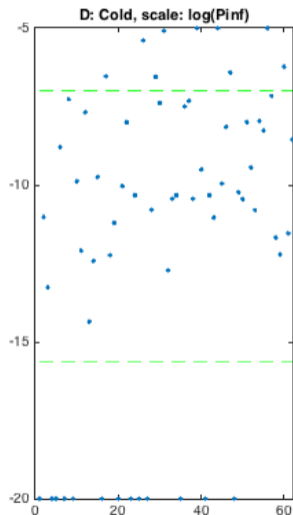
Primal/dual infeasibilities:

Step D: Double MINOS, cold start, scale (repeat)

Scale option 2
 Feasibility tol $1e-7$
 Optimality tol $1e-7$

LU Partial Pivoting
 LU Factor tol 10.0
 LU Update tol 10.0
 Expand freq 100000

$\epsilon = 2.2e-16$



Primal/dual infeasibilities:

Step Q1: Quad MINOS, warm start, scale

Scale option 2

Feasibility tol $1e-15$

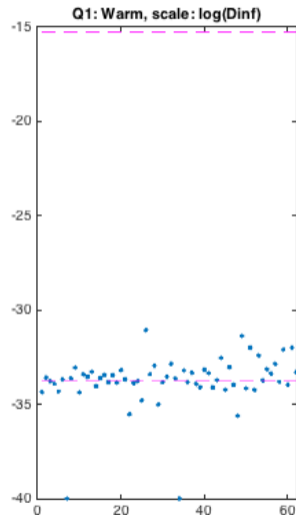
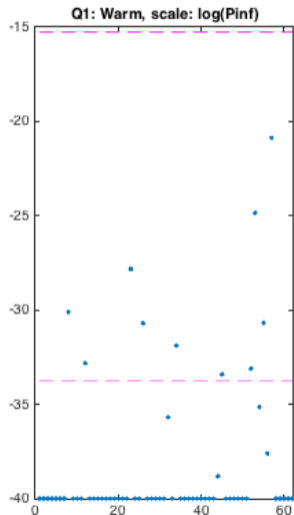
Optimality tol $1e-15$

LU Partial Pivoting

LU Factor tol 10.0

LU Update tol 10.0

$\epsilon = 1.9e-35$



Primal/dual infeasibilities:

Step Q2: Quad MINOS, warm start, no scale

Scale option 0

Feasibility tol $1e-15$

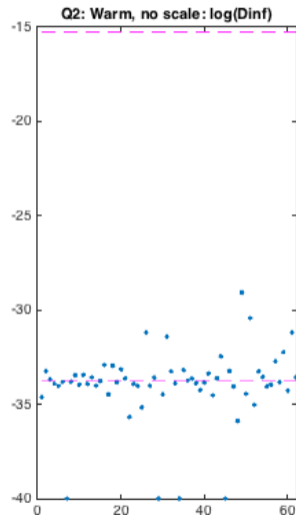
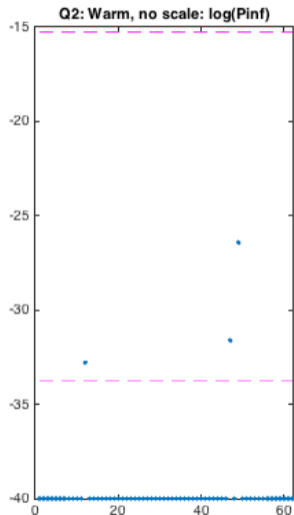
Optimality tol $1e-15$

LU Partial Pivoting

LU Factor tol 5.0

LU Update tol 5.0

$\epsilon = 1.9e-35$



Multiscale NLPs

Systems biology FBA problems with variable μ
Palsson Lab, UC San Diego, 2014–2016

ME models with nonlinear constraints

Coupling constraints can be functions of the organism's growth rate μ .

Lerman et al. (UCSD) consider growth-rate optimization nonlinearly with a single μ as the objective. Nonlinear constraints

$$\frac{v_i}{v_j} \leq \mu \quad \equiv \quad v_i \leq \mu v_j$$

are linear if μ is fixed. Can use a **binary search to find the largest $\mu \in [\mu_{\min}, \mu_{\max}]$ that keeps the associated LP feasible.**

Requires **reliable solution of a sequence of related LPs.**

tinyME (2500 × 2800 nonlinear ME model)

Nonlinear FBA formulation, Laurence Yang, UCSD, Dec 2014

$$\begin{array}{ll}
 \max & \mu \\
 \text{st} & \mu Ax + Bx = 0 \\
 & Sx = b \\
 & \text{bounds on } x
 \end{array}
 \quad \equiv \quad
 \begin{array}{ll}
 \max & \mu \\
 \text{st} & \mu Ax + w = 0 \\
 & Bx - w = 0 \\
 & Sx = b \\
 & \text{bounds on } x, \text{ no bounds on } w
 \end{array}$$

- $\mu = x_1$ First columns of A , B are empty
- Binary search
25 LP subproblems would give 8 digits (really need quad Simplex)
- Instead, apply quad MINOS LCL method = Linearly Constrained Lagrangian
6 NLP subproblems (with linearized constraints) give 20 digits

solveME (11000 \times 19000 nonlinear ME model)

2600 nonlinear constraints, 16000 nonlinear variables

Laurence Yang, UCSD, Sep 2015

Itn 32 -- linear constraints satisfied.

Calling funcon. mu = 0.832815729997476367249118875820191994

Major	minor	step	objective	Feasible	Optimal	nsb	ncon	penalty	BSwap
1	32T	0.0E+00	8.32816E-01	4.3E-13	1.0E+03	0	4	1.0E+02	0
19	40T	1.0E+00	8.32816E-01	2.5E-16	1.0E-03	0	743	1.0E+02	0
20	40T	1.0E+00	8.32816E-01	1.0E-21	9.3E-04	0	784	1.0E+02	0
23	40T	1.0E+00	8.55337E-01	3.4E-07	5.7E-05	0	907	1.0E+02	0
24	40T	1.0E+00	8.55664E-01	2.1E-08	6.6E-07	0	948	1.0E+02	0
25	11	1.0E+00	8.55664E-01	7.0E-17	1.8E-11	0	961	1.0E+02	0
26	0	1.0E+00	8.55664E-01	9.3E-19	8.0E-29	0	962	1.0E+01	0

EXIT -- optimal solution found

ME 2.0 (large FBA and FVA problems)

71,000 \times 80,000 LPs
 Laurence Yang, UCSD, Sep 2015

Quad MINOS cold start: \sim 3 hours

FVA problems: min and max individual variables v_j

Reaction	Protein	Double CPLEX		Quad MINOS
		v_{\min}	v_{\max}	$v_{\min} = v_{\max}$
translation_b0169	RpsB	30.715011	30.712581	30.719225
translation_b0025	RibF	0.212807	0.211712	0.210161
translation_b0071	LeuD	0.303304	0.765585	0.303634
translation_b0072	LeuC	0.303304	0.681146	0.303634

quadQP

Laurence Yang, UCSD, Jul 2016

$$\min c^T x + \frac{1}{2} x^T H x \quad \text{st} \quad Ax + s = b \text{ and bounds,} \quad \begin{array}{l} H \quad 4146 \times 4146 \\ A \quad 12667 \times 21126 \end{array}$$

Itn	rg	ninf	sinf	objective
1	-1.1E+11	159	7.715E+00	0.00000000E+00
500	-1.9E+10	159	7.715E+00	0.00000000E+00
900	-1.3E+14	159	7.715E+00	0.00000000E+00
3000	-1.5E+05	11	9.784E-04	0.00000000E+00
3400	-5.1E-02	4	6.616E-05	0.00000000E+00
Itn	3554	-- feasible solution.		Objective = 7.634331271E-03

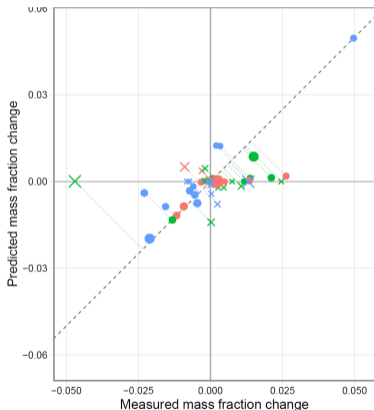
Itn	rg	ninf	sinf	objective	nobj	nsb
4000	-4.8E-02	0	0.000E+00	3.08300094E-03		
5000	-1.9E-04	0	0.000E+00	1.03117165E-03	1406	1
5200	-1.9E-05	0	0.000E+00	1.02662098E-03	1436	2
5300	-4.6E-09	0	0.000E+00	1.02660849E-03	1466	4

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246 secs

Refining ME-computed proteomes



Proteome: the set of all proteins that is or can be expressed by an organism

Mass fraction: relative abundance of a given protein relative to all proteins (i.e., the proteome)

Includes solving quad-precision QP

Itn	rg	ninf	sinf	objective	nobj	nsb
4700	-2.9E-03	0	0.000E+00	1.35572488E-03		
4800	-1.2E-03	0	0.000E+00	1.03964380E-03		
4900	-4.1E-04	0	0.000E+00	1.03667479E-03		

Itn	rg	ninf	sinf	objective	nobj	nsb
5000	-1.9E-04	0	0.000E+00	1.03117165E-03	1406	1
5100	-6.6E-05	0	0.000E+00	1.02674422E-03	1423	1
5200	-1.9E-05	0	0.000E+00	1.02662098E-03	1436	2
5300	-4.6E-09	0	0.000E+00	1.02660849E-03	1466	4

```

Itn      5318 --      25 nonbasics set on bound, basics recomputed
Itn      5318 -- feasible solution. Objective = 1.026608483E-03
  
```

EXIT -- optimal solution found

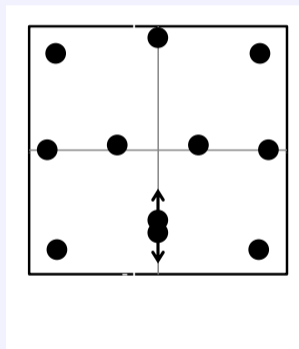
No. of iterations	5318	Objective value	1.0266084830E-03
No. of major iterations	1	Linear objective	0.0000000000E+00
Penalty parameter	100.000000	Nonlinear objective	1.0266084830E-03
No. of calls to funobj	1489	No. of calls to funcon	0
No. of superbasics	2	No. of basic nonlinear	12614
No. of degenerate steps	3686	Percentage	69.31
Max x (scaled)	12400 3.3E-01	Max pi (scaled)	662 1.5E+04
Max x	10269 1.9E+01	Max pi	6083 3.4E+02
Max Prim inf(scaled)	7860 1.1E-25	Max Dual inf(scaled)	9618 1.5E-21
Max Primal infeas	8602 1.7E-22	Max Dual infeas	9618 4.8E-25

Quasi-Newton optimization with finite-difference gradients

Design of computer experiments

Selden Crary, indie-physicist, 2015

$n = 11$ points (x_i, y_i) on $[-1, 1]$ square (one twin-point)



$[d,n,p,\theta_1,\theta_2]=[2,11,2,0.128,0.069]$

Design of computer experiments

Selden Crary, indie-physicist, 2015

IMSPE-optimal designs (integrated mean-squared prediction error)

$$\min 1 - \text{trace}(L^{-1}R)$$

L and R : symmetric matrices of order $n + 1$
with L increasingly ill-conditioned if points approach each other

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with L increasingly ill-conditioned if points approach each other

2D, Gaussian covariance parameters $\sigma, \theta_1, \theta_2$

$$L = \begin{bmatrix} 0 & \mathbf{1}^T \\ \mathbf{1} & V \end{bmatrix}, \quad R \propto \int_{-1}^1 \int_{-1}^1 \begin{bmatrix} 1 \\ v \end{bmatrix} [1 \quad v^T] dx dy$$

$$V_{ij} = \sigma^2 e^{-\theta_1(x_i - x_j)^2 - \theta_2(y_i - y_j)^2}, \quad v_i \text{ functions of } \exp(\cdot) \text{ and } \text{erf}(\cdot)$$

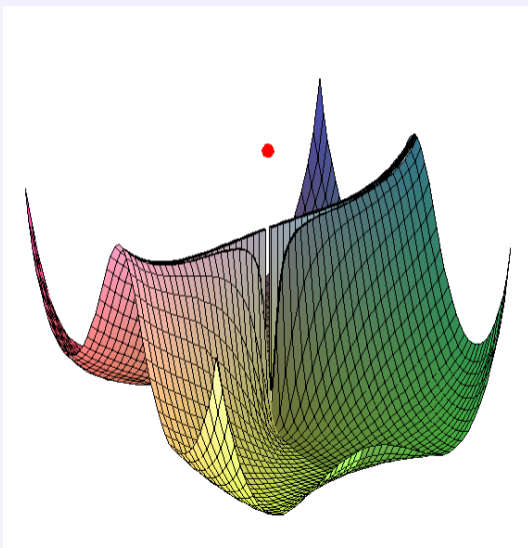
Pagoda plot

Selden Crary, 2015

 C^∞ a.e.

"Post", not pole

Need multistarts



With Maple, Selden has found twin-points, triple-points, ... and a new class of rational functions (the Nu class)

Smooth sailing through the black-hole/white-hole event horizon

No string theory

No complex numbers in quantum mechanics

...

IMSPE, 2D, $n = 11$, $\theta_1 = 0.128$, $\theta_2 = 0.069$

Selden Crary

Quad MINOS unconstrained optimization $\in \mathbb{R}^{22}$ without gradients 6 secs

Itn	ph	pp	rg	step	objective	nobj	nsb	cond(H)
1	4	0	3.9E-05	1.0E+03	2.47305090E-05	57	22	4.5E+01
50	4	1	6.4E-07	6.2E+00	6.01181966E-06	1384	22	2.1E+04
100	3	1	-7.6E-08	1.1E+00	5.65611811E-06	2726	22	1.4E+05
150	4	1	8.5E-07	6.0E+00	5.11053080E-06	4102	22	8.2E+03
200	4	1	2.6E-08	1.1E-01	5.02762155E-06	5464	22	1.0E+07
239	4	1	1.1E-07	1.5E-06	5.02762154E-06	7478	22	1.0E+10

alfmax = 3.1E+04 pnorm = 1.5E-04 gnorm = 1.1E-07 g'p = -2.6E-15 numf = 15
 Search exit 7 -- too many functions.

EXIT -- optimal solution found

No. of iterations	239	Objective value	5.0276215358E-06
No. of calls to funobj	7538	Calls with mode=2 (f, known g)	244
Calls for forward differencing	4466	Calls for central differencing	1716
Max Primal infeas	0 0.0E+00	Max Dual infeas	2 1.1E-07
			(usually 1.0E-15)

IMSPE linear algebra question

L, R real, symmetric, indefinite, ill-conditioned

$$\min 1 - \text{trace}(L^{-1}R)$$

cf. GEV problem $Rv = \lambda Lv$

$$\text{trace}(L^{-1}R) = \sum \lambda_i$$

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- Will QZ compute real λ_i ?

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Yuji Nakatsukasa was in Tokyo; developing `qdwhep.m` for $Ax = \lambda Bx$ (real, symmetric)

- Congruence transformations are real
- Eigenvalues can be complex conjugate pairs
- $\text{trace}(B^{-1}A) = \sum \lambda_i$ will be real

Conclusions

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- Iterative refinement of $Bp = a$, $B^T y = c$ is a sparing use of Quad, but doesn't help $B = LU$ (with **basis repair**).
Quad everywhere is the humane approach!

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Quad everywhere is the humane approach!
- Same difficulty with SoPlex80bit + refinement with rational arithmetic.
Quad SoPlex is in the making!

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- Philip Gill, Elizabeth Wong (UCSD)
- Ed Klotz (IBM CPLEX), Yuekai Sun (ICME), Jon Dattorro (Stanford)
- Selden Crary (Palo Alto), Richard Martinez (MS&E)

Imperial Palace, Tokyo



Tokyo subway



Welcome home



FAQ

- Is quadMINOS available? Yes, in the openCOBRA toolbox
<http://opencobra.github.io/cobratoolbox/>
- Can quadMINOS be called from Matlab or Tomlab? Yes via system call (not Mex)
- Is quadMINOS available in GAMS? Yes for LPs
- How about AMPL? No, but should be feasible
- Is there a quadSNOPT? Yes, in f90 SNOPT9 we change 1 line
- Can CPLEX / Gurobi / Mosek / ... help? Yes, they can provide Presolve and Warm start, especially from GAMS
- Will Quad hardware eventually be standard? We wish, but Kahan is pessimistic