BIOSTAT M280/BIOMATH 280/STAT M230: Statistical Computing

Tue/Thu 1:00pm-2:50pm, CHS 51-279 Instructor: Dr. Hua Zhou, huazhou@ucla.edu

1 Lecture 1: Jan 5

Today

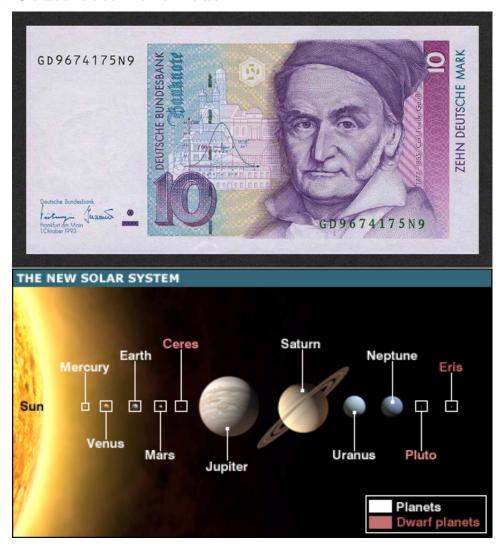
- Introduction and course logistics
- Computer storage and arithmetic
- Homework (not graded): fill out a short survey at https://www.surveymonkey. com/r/G8BCVVM
- Homework (not graded): Read the Introduction and Foundations of *Advanced R* by Hadley Wickham http://adv-r.had.co.nz. Install R. Try examples while you read the book. Do the quizzes.

What is statistics?

- People collect data in order to answer certain questions. (Bio)statisticians's job is to help make sense of data.
- Statistics, the science of *data analysis*, is the applied mathematics in the 21st century.
- Read papers *The future of data analysis* by John Tukey (http://hua-zhou.github.io/teaching/biostatm280-2016winter/readings/Tukey61FutureDataAnalysis.pdf) and *50 years of data sicence* by David Donoho (http://hua-zhou.github.

io/teaching/biostatm280-2016winter/readings/Donoho15FiftyYearsDataScience.pdf).

How Gauss became famous?



- 1801, Dr. Carl Friedrich Gauss, 24; proved Fundamental Theorem of Algebra; wrote the book Disquisitiones Arithmetic, which is still being studied today.
- 1801, Jan 1-Feb 11 (41 days), astronomer Piazzi observed Ceres (a dwarf planet), which was then lost behind sun.
- 1801, Aug—Sep, futile search by top astronomers; Laplace claimed it unsolvable.

- 1801, Oct-Nov, Gauss did calculations by method of least squares.
- 1801, Dec 31, astronomer von Zach re-located Ceres according to Gauss' calculation.
- 1802, Summarische Übersicht der Bestimmung der Bahnen der beiden neuen Hauptplaneten angewandten Methoden, considered the origin of linear algebra.
- 1807, Professor of Astronomy and (the first) Director of Göttingen Observatory in remainder of his life.
- 1809, Theoria motus corporum coelestium in sectionibus conicis solum ambientium (Theory of motion of the celestial bodies moving in conic sections around the Sun); birth of the Gaussian (normal) distribution, as an attempt to rationalize the method of least squares.
- 1810, Laplace consolidated the importance of Gaussian distribution by proving the central limit theorem.
- 1829, Gauss-Markov Theorem. Under Gaussian error assumption (actually only uncorrelated and homoscedastic needed), least square solution is the best linear unbiased estimate (BLUE), i.e., it has the smallest variance and thus MSE among all linear unbiased estimators. Note other estimators such as the James-Stein estimator may have smaller MSE, but they are *nonlinear*.

For more details of the story

- http://www.keplersdiscovery.com/Asteroid.html
- Teets and Whitehead (1999)

ARTICLES

The Discovery of Ceres: How Gauss Became Famous

DONALD TEETS KAREN WHITEHEAD South Dakota School of Mines and Technology Rapid City, SD 57701

"The Duke of Brunswick has discovered more in his country than a planet: a super-terrestrial spirit in a human body."

These words, attributed to Laplace in 1801, refer to the accomplishment of Carl Friedrich Gauss in computing the orbit of the newly discovered planetoid *Ceres Ferdinandea* from extremely limited data. Indeed, although Gauss had already achieved some fame among mathematicians, it was his work on the Ceres orbit that "made Gauss a European celebrity—this a consequence of the popular appeal which astronomy has always enjoyed..." [2]. The story of Gauss's work on this problem is a good one and is often told in biographical sketches of Gauss (e.g., [2], [3], [6]), but the mathematical details of how he solved the problem are invariably omitted from such historical works. We are left to wonder how did he do it? *Inst. how. did Gauss*

• Stigler (1986) gives a more comprehensive account of the origin of the method of least squares.

Gauss' story

- Motivated by a real problem.
- Heuristic solution: method of least squares.
- Solution readily verifiable: Ceres was re-discovered!
- Algorithmic development: linear algebra, Gaussian elimination, FFT (fast Fourier transform).
- Theoretical justification: Gaussian distribution, Gauss-Markov theorem.

A sampler by Marc Coram

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to bat-rb. con todo mi respeto. i was sitting down playing chess with danny de emf and boxer de el centro was sitting next to us. boxer was making loud and loud voices so i tell him por favor can you kick back bonie cause im playing chess a minute later the vato starts back up again so this time i tell him con respecto homie can you kick back. the vato stop for a minute and he starts up again so i tell him check this out shut the foot up cause im tired of your voice and if you got a problem with it we can go to celda and handle it. i really felt disrespected thats why i told him. anyways after i tell him that the next thing I know that vato slashes me and leaves. dy the time i figure im hit i try to get away but the c.o. is walking in my direction and he gets me right dy a celda. so go to the hole. when im in the hole my home boys hit doxer so now "b" is also in the hole. while im in the hole im getting schoold wrong and

- A consulting project by Marc Coram (then a graduate student in statistics at Stanford); customer is a professor in political science, who wants to understand a cryptic message circulated in a state prison.
- Marc modeled letter sequence by a Markov chain $(26 \times 26 \text{ transition matrix})$ and estimated transition probabilities from War and Peace.

- Now each mapping σ yields a likelihood $f(\sigma)$ of the symbol sequence.
- Find the σ that maximizes f. Sample space is at least $26! = 4.0329 \times 10^{26}$. Combinatorial optimization hard!
- Metropolis algorithm: At each iteration, generate a new σ' by random transposition of two letters; accept σ' with probability min $\left\{\frac{f(\sigma')}{f(\sigma)}, 1\right\}$.

Marc Coram's story

- Motivated by a real problem.
- Solution readily verifiable: we can read it!
- Algorithm development: Metropolis sampler is one of top 10 algorithms in the 20th century.
- Read Diaconis (2009) for more details.

What is this course about?

- Not a course on "packages and languages for data analysis". It does not answer questions such as "How to fit a linear mixed model in SAS, SPSS or R?"
- Not a programming course, although programming is *extremely* important and we do homework in R.
- This course is about "numerical methods in statistics". Our focus is on algorithms.

The form of a mathematical expression and the way the expression should be evaluated in actual practice may be quite different.

James Gentle

For a common numerical task in statistics, say the least squares solution $\widehat{\boldsymbol{\beta}} = (\boldsymbol{X}^{\mathsf{T}}\boldsymbol{X})^{-1}\boldsymbol{X}^{\mathsf{T}}\boldsymbol{y}$, we need to know which methods/algorithms are out there and what are their advantages and disadvantages. You will *fail* this course if you use

solve(t(X) %*% X) %*% t(X) %*% y

Using lm(y ~ X) is correct (and efficient) but is not the purpose of this course. We want to understand what is going on when calling the lm() function.

Algorithms for the Ages

"Great algorithms are the poetry of computation," says Francis Sullivan of the Institute for Defense Analyses' Center for Computing Sciences in Bowie, Maryland. He and Jack Dongarra of the University of Tennessee and Oak Ridge National Laboratory have put together a sampling that might have made Robert Frost beam with pride—had the poet been a computer jock. Their list of 10 algorithms having "the greatest influence on the development and practice of science and engineering in the 20th century" appears in the January/February issue of *Computing in Science & Engineering*. If you use a computer, some of these algorithms are no doubt crunching your data as you read this. The drum roll, please:

1946: The Metropolis Algorithm for Monte Carlo. Through the use of random processes, this algorithm offers an efficient way to stumble toward answers to problems that are too complicated to solve exactly.

1947: Simplex Method for Linear Programming. An elegant solution to a common problem in planning and decision-making.

1950: Krylov Subspace Iteration Method. A technique for rapidly solving the linear equations that abound in scientific computation.

1951: The Decompositional Approach to Matrix Computations. A suite of techniques for numerical linear algebra.

1957: The Fortran Optimizing Compiler. Turns high-level code into efficient computer-readable code.

1959: QR Algorithm for Computing Eigenvalues. Another crucial matrix operation made swift and practical.

1962: Quicksort Algorithms for Sorting. For the efficient handling of large databases.

1965: Fast Fourier Transform. Perhaps the most ubiquitous algorithm in use today, it breaks down waveforms (like sound) into periodic components.

1977: Integer Relation Detection. A fast method for spotting simple equations satisfied by collections of seemingly unrelated numbers.

1987: Fast Multipole Method. A breakthrough in dealing with the complexity of n-body calculations, applied in problems ranging from celestial mechanics to protein folding.

Syllabus

Check course website frequently for updates and announcements. http://hua-zhou.github.io/teaching/biostatm280-2016winter Lecture notes will be updated and posted after each lecture.

Computer storage and arithmetic

Elementary units of computer storage:

- bit = "binary" + "digit" (coined by statistician John Tukey).
- byte = 8 bits.
- $kB = kilobyte = 10^3$ bytes.
- $MB = megabytes = 10^6 bytes.$
- $GB = gigabytes = 10^9 bytes.$
- TB = terabytes = 10^{12} bytes.
- PB = petabytes = 10^{15} bytes.

Storage of characters

ASCII control characters				ASCII printable characters						Extended ASCII characters							
00	NULL	(Null character)	32	space	64	@	96	٠,	1	28	Ç	160	á	192	L	224	Ó
01	SOH	(Start of Header)	33		65	Α	97	а	1	29	ü	161	ĺ	193	Τ.	225	ß
02	STX	(Start of Text)	34		66	В	98	b	1	30	é	162	Ó	194	т	226	Ô
03	ETX	(End of Text)	35	#	67	С	99	С	1	131	â	163	ú	195	-	227	Ò
04	EOT	(End of Trans.)	36	\$	68	D	100	d	1	32	ä	164	ñ	196	-	228	Õ
05	ENQ	(Enquiry)	37	%	69	E	101	е	1	33	à	165	Ñ	197	+	229	Õ
06	ACK	(Acknowledgement)	38	&	70	F	102	f	1	34	å	166	a	198	ä	230	μ
07	BEL	(Bell)	39		71	G	103	g	1	35	ç	167	0	199	Ã	231	þ
08	BS	(Backspace)	40	(72	Н	104	h		36	ê	168	ż	200	L	232	Þ
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10	LF	(Line feed)	42	*	74	J	106	j	1	38	è	170	7	202	T	234	Û
11	VT	(Vertical Tab)	43	+	75	K	107	k	1	39	ï	171	1/2	203	TE	235	Ù
12	FF	(Form feed)	44	,	76	L	108	- 1	1	40	î	172	1/4	204	ŀ	236	Ý
13	CR	(Carriage return)	45		77	M	109	m	1	41	ì	173	i	205	=	237	Ý
14	SO	(Shift Out)	46		78	N	110	n	1	42	Ä	174	«	206	#	238	_
15	SI	(Shift In)	47	1	79	0	111	0	1	43	Α	175	»	207	n	239	
16	DLE	(Data link escape)	48	0	80	Р	112	р	1	44	É	176		208	õ	240	=
17	DC1	(Device control 1)	49	1	81	Q	113	q	1	45	æ	177	- 8	209	Ð	241	±
18	DC2	(Device control 2)	50	2	82	R	114	r	1	46	Æ	178		210	Ê	242	_
19	DC3	(Device control 3)	51	3	83	S	115	s	1	47	ô	179	T	211	Ë	243	3/4
20	DC4	(Device control 4)	52	4	84	T	116	t	1	48	ö	180	+	212	È	244	1
21	NAK	(Negative acknowl.)	53	5	85	U	117	u	1	49	ò	181	À	213	- 1	245	§
22	SYN	(Synchronous idle)	54	6	86	V	118	V	1	150	û	182	Â	214	ĺ	246	÷
23	ETB	(End of trans. block)	55	7	87	W	119	w	1	151	ù	183	À	215	Î	247	
24	CAN	(Cancel)	56	8	88	Х	120	х	1	52	ÿ	184	©	216	Ϊ	248	
25	EM	(End of medium)	57	9	89	Υ	121	У	1	53	Ö	185	4	217	7	249	-
26	SUB	(Substitute)	58	:	90	Z	122	z	1	54	Ü	186		218	Г	250	
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30	RS	(Record separator)	62	>	94	^	126	~	1	58	×	190	¥	222	ĺ	254	
31	US	(Unit separator)	63	?	95	_			1	59	f	191	7	223		255	nbsp
127	DEL	(Delete)				_											

- ullet Plain text files are stored in the form of characters: .r, .c, .cpp, .tex, .html,
- ASCII (American Code for Information Interchange): 7 bits, only $2^7 = 128$ characters, "Hua" corresponds to "48 75 61 (Hex) = 72 117 97 (Dec) = 1001000 1110101 1100001".

- Extended ASCII: 8 bits, $2^8 = 256$ characters.
- Unicode: UTF-8, UTF-16 and UTF-32 support many more characters including foreign characters; last 7 digits conform to ASCII. UTF-8 is the current dominant character encoding on internet.

2 Lecture 2, Jan 7

Announcements

- Location change: class meeting is changed to CHS 51-279, effective Jan 7.
- Enrollment cap is raised to 38 (open).
- Use RStudio for R programming.

Last time

- Introduction. Gauss (least squares to find Ceres)—optimization, Marc Coram (decipher a note circulating in jail)—sampling. Two major modes of statistical computing.
- Course content, logistics.
- Computer representation of characters (ASCII, unicode) and integers (fixed point number system).

Today

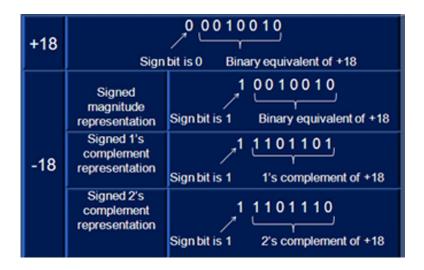
- Computer representation and arithmetic of integers: fixed-point numbers.
- Computer representation and arithmetic of real numbers: floating-point numbers.

Fixed-point number system

Fixed-point number system \mathbb{I} is a computer model for integers \mathbb{Z} . One storage unit may be M = 8/16/32/64 bit.

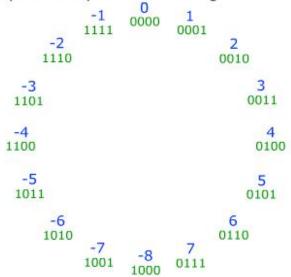
- The number of bits and method of representing negative numbers vary from system to system. The integer type in R has M = 32 bits. MATLAB has (u)int8, (u)int16, (u)int32, (u)int64. JULIA has even more choices such as Int128, UInt128, BigInt, ...
- First bit indicates sign: 0 for nonnegative numbers, 1 for negative numbers.

• "two's complement representation" for negative numbers. (i) Sign bit is set to 1, (ii) remaining bits are set to opposite values, (iii) 1 is added to the result.



- Range of representable integers by M-bit storage is $[-2^{M-1}, 2^{M-1} 1]$ (don't need to represent 0 anymore so *could* have capacity for 2^{M-1} negative numbers).
- For M = 8, [-128, 127]. For M = 16, [-65536, 65535]. For M = 32, [-2147483648, 2147483647].
- The smallest representable integer in R is $-2^{31} + 1 = -2147483647$.
- For unsigned integers such as in MATLAB, the range is $[0, 2^M 1]$.

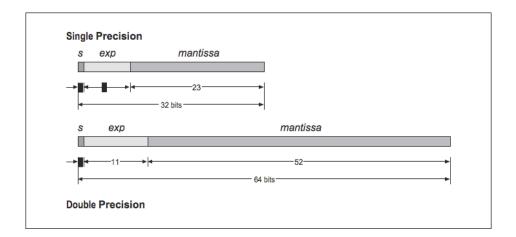
Two's Complement representation using 4 bit binary strings



- An integer -i in the interval $[-2^{M-1}, -1]$ would be represented by the same bit pattern by which the nonnegative integer $2^M i$ is represented, treating the sign bit as a regular numeric bit. For example, (-18) = 11101110, $(2^8) (18) = (238) = 11101110$. Two's complement is subtracting the nonnegative integer i from $1 \dots 1 + 1 = 10 \dots 0 = (2^M)$.
- Addition and subtraction are much simpler in two's complement representation. Why? It respects modular arithmetic nicely (look at the diagram). E.g., (3)+(4) = 0011 + 0100 = 0111 = (7), (3)+(-5) = 0011 + 1011 = 1110 = (-2), (3) + (7) = 0011 + 0111 = 1010 = (-6) (overflow), (-3) + (-7) = 1101 + 1001 = 0110 = (6) (underflow).

- Keep track of overflow and underflow. If the result of a summation is R, which must be in the set $[-2^{M-1}, 2^{M-1} + 1]$, there are only three possibilities for the true sum: R, $R + 2^M$ (overflow), or $R 2^M$ (underflow).
 - When adding two nonnegative integers: 0XX...X + 0YY...Y, where X and Y are arbitrary binary digits, we only need to keep track of overflow in this case. If the resulting binary number has a leading bit of 1, we know overflow occurs since the sum cannot be negative. We should treat that sign bit as a regular numeric bit. In other words, we crossed the upper boundary 100 and should add 2^M to the result. If the resulting binary number has a leading bit of 0, no overflow occurs.
 - When adding two negative integers: 1XX...X + 1YY...Y, where X and Y are arbitrary binary digits, we only need to keep track of underflow in this case. If the resulting binary number has a leading bit of 0, we know underflow occurs since the sum cannot be positive. We crossed the lower boundary 000 and should subtract 2^M from the result. If the resulting binary number has a leading bit of 1, no underflow occurs.
 - When adding a negative integer and a nonnegative integer: 1XX...X + 0YY...Y, the result is always between the two summands. No overflow or underflow could happen.
- R reports NA for integer overflow and underflow. Other languages, e.g., Julia simply outputs the result according modular arithmetic.

Floating-point number system



Floating-point number system \mathbb{F} is a computer model for real numbers \mathbb{R} .

- A real number is represented by $\pm d_0.d_1d_2\cdots d_p\times b^e$ (scientific notation).
- Parameters for a floating-point number system: base (or radix), range of fraction (or mantissa, significand), range of exponent.
- Non-uniqueness of representation. normalized/denormalized significant digits. E.g., $+18 = +1.0010 \times 2^4 \text{(normalied)} = +0.10010 \times 2^5 \text{(denormalized)}.$
- Bias (or excess): actual exponent is obtained by subtracting bias from the value of exponent evaluated regardless of sign digit.
- IEEE 754.
 - Single precision (32 bit): base 2, p=23 (23 significant bits), $e_{\rm max}=127$, $e_{\rm min}=-126$ (8 exponent bits), bias=127. $e_{\rm min}-1$ and $e_{\rm max}+1$ are reserved for special numbers. This implies a maximum magnitude of $\log_{10}(2^{127})\approx 38$ and precision to $\log_{10}(2^{23})\approx 7$ decimal point. $\pm 10^{\pm 38}$.
 - Double precision (64 bit): base 2, p=52 (52 significant bits), $e_{\rm max}=1023$, $e_{\rm min}=-1022$ (11 exponent bits), bias=1023. This implies a maximum magnitude of $\log_{10}(2^{1023})\approx 308$ and precision to $\log_{10}(2^{52})\approx 16$ decimal point. $\pm 10^{\pm 308}$.

• " $(+18) = (2^4 + 2^1) = +1.0010 \times 2^4$ in single precision (0)(10000011)(001000000000000000000).

First is sign bit. Next 8 bits are exponent 131 in ordinary base 2 with a bias of 127. Remaining 23 bits represent the fraction beyond the leading bit, known to be 1. In summary it represents (+18) as $+1.0010 \times 2^4$ in the binary format. (-18) is represented by the same bits except changing the sign bit to 1.

- Special floating-point numbers:
 - Exponent $e_{\text{max}} + 1$ plus a mantissa of 0 means $\pm \infty$.
 - Exponent $e_{\text{max}} + 1$ plus a nonzero mantissa means NaN. NaN could be produced from 0 / 0, 0 * Inf, ... In general $NaN \neq NaN$ bitwise.
 - Exponent e_{\min} 1 with a mantissa of all 0s represents the real number 0.
 - Exponent $e_{\min} 1$ with a nonzero mantissa are for numbers less than $b^{e_{\min}}$. Numbers are de-normalized in the range $(0, b^{e_{\min}})$ – "graceful underflow".
- \mathbb{F} is not a subset of \mathbb{R} , although $\mathbb{I} \subset \mathbb{Z}$.
- For the history of establishment of IEEE-754 standard, see an interview with William Kahan.

http://www.cs.berkeley.edu/~wkahan/ieee754status/754story.html

To summarize

- Single precision: range $\pm 10^{\pm 38}$ with precision up to 7 decimal digits.
- Double precision: range $\pm 10^{\pm 308}$ with precision up to 16 decimal digits.
- The floating-point numbers do not occur uniformly over the real number line.



Fig. 2.4. The Floating-Point Number Line, Nonnegative Half

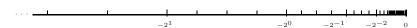


Fig. 2.5. The Floating-Point Number Line, Nonpositive Half



Fig. 2.6. The Floating-Point Number Line, Nonnegative Half; Another View

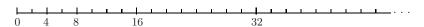


Fig. 2.7. The Fixed-Point Number Line, Nonnegative Half

• Machine epsilons are the spacings of numbers around 1. $\epsilon_{\min} = b^{-p}$ and $\epsilon_{\max} = b^{1-p}$.



Fig. 2.8. Relative Spacings at 1: "Machine Epsilons"

- The variable .Machine in R contains numerical characteristics of the machine.
- How to test inf and nan? In R, is.nan(), is.finite(), is.infinite(). In MATLAB, isinf(), isnan(). In JULIA, isinf(), isnan().

Integers and floating-point numbers in Julia

• R only uses double (64-bit) and 32-bit integer. It can be a downside when dealing with big data.

• Julia offers a rich collection of primitive data types. Read (the excellent) documentation. http://docs.julialang.org/en/release-0.4/manual/integers-and-floating-Notable things include: Int128, UInt128, half precision numbers Float16, arbitrary precision numbers BigInt and BigFloat, rational numbers, ...

Consequences of computer storage/arithmetic

- Be memory conscious when dealing with big data. E.g., human genome has about 3×10^9 bases, each of which belongs to $\{A, C, T, G\}$. How much storage if we store 10^6 SNPs (single nucleotide polymorphisms) of 1000 individuals (1000 Genome Project) as characters (1GB), single (4GB), double (8GB), int64(8GB), int32 (4GB), int16 (2GB), int8 (1GB), PLINK binary format 2bit/SNP (250MB)?
- Know the limit. Overflow and Underflow. For double precision, $\pm 10^{\pm 308}$. In most situations, underflow is preferred over overflow. Overflow often causes crashes. Underflow yields zeros. E.g., in logistic regression, $p_i = \frac{\exp(\boldsymbol{x}_i^\mathsf{T}\boldsymbol{\beta})}{1+\exp(\boldsymbol{x}_i^\mathsf{T}\boldsymbol{\beta})} = \frac{1}{1+\exp(-\boldsymbol{x}_i^\mathsf{T}\boldsymbol{\beta})}$. The former expression can easily lead to $\infty/\infty = NaN$, while the latter expression leads to graceful underflow.
- Be aware of non-uniform distribution of floating-point numbers, in contrast to fixed-point numbers. There are the same number of floating-point numbers in $[b^i, b^{i+1}]$ and $[b^{i+1}, b^{i+2}]$ for $e_{\min} \leq i \leq e_{\max} 2$. It is more dense when closer to zero.
- "Catastrophic cancellation 1". Addition or subtraction of two numbers of widely different magnitudes: a+b or a-b where $a\gg b$ or $a\ll b$. We loose the precision in the number of smaller magnitude. Consider $a=x.xxx...\times 2^0$ and $b=y.yyy...\times 2^{-53}$. What happens when computer calculates a+b? We get a+b=a!
- Another example: What happens when compute $\sum_{x=1}^{\infty} x$ in order? Will the partial sum reach Inf? "A divergent series converges."
- Always try to add numbers of similar magnitude. Rule 1: add small numbers together before adding larger ones. Rule 2: add numbers of like magnitude together (paring). When all numbers are of same sign and similar magnitude, add in pairs so each stage the summands are of similar magnitude.

- E.g., evaluating e^{-20} by Taylor series

$$e^{-x} = 1 - x + x^2/2! - x^3/3! + \cdots$$

gets 6.138e-09, while the true value is about 2.061e-09. Many cancellations accumulate. Anyway, it's not the way to evaluate e^x !

• Sometimes catastrophic cancellation can be avoided. Roots of the quadratic function $ax^2 + bx + c$ are

$$x = \frac{-b \pm \sqrt{b^2 - 4ac}}{2a}.$$

When one root is close to 0, cancellation can happen. We may evaluate one of the root (away from 0) by the formula and then compute the other by relationship $x_1x_2 = c/a$.

• Reading: What every computer scientist should know about floating-point arithmetic by David Goldberg.

http://hua-zhou.github.io/teaching/biostatm280-2016winter/readings/Goldberg91FloatingPoint.pdf

3 Lecture 3, Jan 12

Announcements

- HW1 posted. Due Thu Jan 21 @ 11:59PM. http://hua-zhou.github.io/teaching/biostatm280-2016winter/biostat_m280_2016_hw1.pdf
- Quiz 1 Thu Jan 21 in class.
- Teaching assistant (TA): Max Tolkoff mtolkoff@ucla.edu, office hours M @ 11A-12P, W @ 2P-3P, at the common area of the 6th floor of Gonda.

Last time

- Computer arithmetics: fixed-point numbers and floating-point numbers.
- Consequences of computer arithmetics: range and precision of single/double precision numbers, cancellations, ...

Today

- Computer languages.
- Comparison of R, Matlab, and Julia.
- R, RStudio, RMarkdown, version control.

Computer Languages

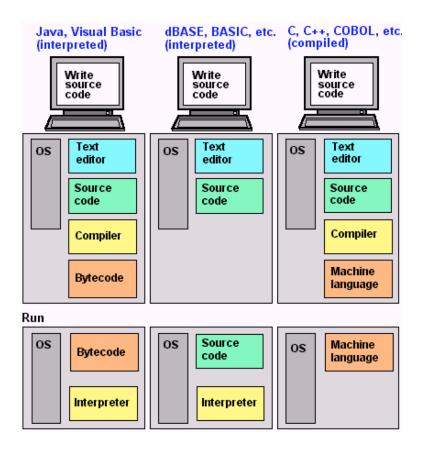
工欲善其事,必先利其器

To do a good job, an artisan needs the best tools.

The Analects by Confucius (about 500 BC)

- What features are we looking for in a language?
 - Efficiency (in both run time and memory) for handling big data.
 - IDE support (debugging, profiling).

- Open source.
- Legacy code.
- Tools for generating dynamic report (reproducibility).
- Adaptivity to hardware evolution (parallel and distributed computing).



• Types of languages

- 1. Compiled languages: C/C++, FORTRAN, ...
 - Directly compiled to machine code that is executed by CPU
 - Pros: fast, memory efficient
 - Cons: longer development time, hard to debug
- 2. Interpreted language: R, MATLAB, SAS IML, JAVASCRIPT, BASIC, ...
 - Interpreted by interpreter
 - Pros: fast prototyping

- Cons: excruciatingly slow for loops
- 3. Mixed (dynamic) languages: Julia, Python, JAVA, Matlab (JIT), R (JIT),
 - Compiled to bytecode and then interpreted by virtual machine
 - Pros: relatively short development time, cross-platform, good at data preprocessing and manipulation, rich libraries and modules
 - Cons: not as fast as compiled language
- 4. Script languages: shell scripts, Perl, ...
 - Extremely useful for data preprocessing and manipulation
 - E.g., massage the Yelp (http://www.yelp.com/dataset_challenge) data before analysis

```
hzhou3@zhou-lap2:yelp_phoenix_academic_dataset $ ls -all
     total 452432
  drwx------@ 9 hzhou3 staff
drwxr-xr-x 7 hzhou3 staff
                                                                                                                                                                                                        306 Mar 28 10:55 ...
                                                                                                                                                                                      300 Mar 26 10:35 ./
238 Apr 10 09:10 ../
6148 Mar 28 10:55 .DS_Store
140579 Mar 26 18:44 READ_FIRST-Phoenix_Academic_Dataset_Agreement-3-11-13.pdf
421 Mar 26 17:10 notes.txt
     -rw-r--r--@ 1 hzhou3
                                                                                                                              staff
      -rw-r--r--@ 1 hzhou3
                                                                     1 hzhou3
                                                                                                                             staff
                                                                                                                             staff 421 Mar 26 17:10 notes.txt
staff 4287324 Mar 28 13:19 yelp_academic_dataset_business.json
staff 3519126 Mar 26 17:54 yelp_academic_dataset_checkin.json
staff 216292386 Mar 28 13:30 yelp_academic_dataset_review.json
staff 7374045 Mar 26 17:54 yelp_academic_dataset_user.json
      -rw-r--r-@ 1 hzhou3
                                                                                hzhou3
       -rw-r--r--@ 1 hzhou3
-rw-r--r-@ 1 hzhou3 staff 216292386 Mar 28 13:30 yelp_academic_dataset_review.json
hzhou3@zhou-lap2:yelp_phoenix_academic_dataset $ head yelp_academic_dataset_user.json
hzhou3@zhou-lap2:yelp_phoenix_academic_dataset $ head yelp_academic_dataset_user.json
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                                                                     1 hzhou3
```

5. Database languages: SQL, Hadoop. Data analysis never happens if we do not know how to retrieve data from databases.

Messages

 To be versatile in the big data era, be proficient in at least one language in each category.

- To improve efficiency of interpreted languages such as R or MATLAB, avoid loops as much as possible. Aka, vectorize code "The only loop you are allowed to have is that for an iterative algorithm." For some tasks where looping is necessary, consider coding in C or For-TRAN. It is "convenient" to incorporate compiled code into R or MATLAB. But do this **only after profiling!**
 - Success stories: glmnet and lars packages in R are based on FORTRAN.
- Modern languages such as Julia are trying to bridge the gap between interpreted and compiled languages. That is to achieve efficiency without vectorizing code.
- Language features of R, Matlab, and Julia:

Features	R	Matlab	JULIA		
Open source	©	(3)	©		
IDE	RStudio ©©	000	②		
Dynamic document	$\odot \odot \odot (RMarkdown)$	000	$\odot \odot (\mathrm{IJulia})$		
Multi-threading	${\tt parallel} \ {\rm pkg}$	©	©		
JIT	${ t compiler} \ { t pkg}$	☺	©		
Call C/Fortran	wrapper, Rcpp	wrapper	no glue code		
Call shared library	wrapper	wrapper	no glue code		
Typing	②	© ©	000		
Pass by reference	②	©	000		
Linear algebra	②	MKL, Arpack	OpenBLAS, Eigpack		
Distributed computing	②	©	000		
Sparse linear algebra	② (Matrix package)	000	000		
Documentation	②	000	© ⊕		
Profiler	©	000	©©©		

• Benchmark code R-benchmark-25.R from http://r.research.att.com/benchmarks/ R-benchmark-25.R covers many commonly used numerical operations used in statistics. We ported (literally) to MATLAB and Julia and report the run times (averaged over 5 runs) here.

Matlab benchmark code:

http://hua-zhou.github.io/teaching/biostatm280-2016winter/benchmark_matlab.m

Julia benchmark code:

http://hua-zhou.github.io/teaching/biostatm280-2016winter/benchmark_julia.jl

Machine specs: Intel i7 @ 2.6GHz (4 physical cores, 8 threads), 16G RAM, Mac OS 10.11.2.

Test	R 3.2.2	Matlab R2014a	JULIA 0.4.2
Matrix creation, trans, deformation (2500×2500)	0.77	0.17	0.14
Power of matrix $(2500 \times 2500, A.^{1000})$	0.26	0.11	0.21
Quick sort $(n = 7 \times 10^6)$	0.76	0.24	0.69
Cross product (2800 × 2800, $A^T A$)	10.72	0.35	0.31
LS solution $(n = p = 2000)$	1.28	0.07	0.09
FFT $(n = 2, 400, 000)$	0.40	0.04	0.64
Eigen-values (600×600)	0.82	0.31	0.57
Determinant (2500×2500)	3.76	0.18	0.17
Cholesky (3000×3000)	4.23	0.15	0.20
Matrix inverse (1600×1600)	3.37	0.16	0.21
Fibonacci (vector calc)	0.34	0.17	0.68
Hilbert (matrix calc)	0.21	0.07	0.01
GCD (recursion)	0.31	0.14	0.12
Toeplitz matrix (loops)	0.36	0.0014	0.02
Escoufiers (mixed)	0.45	0.40	0.21

• A slightly more complicated (or realistic) example taken from Doug Bates's slides http://www.stat.wisc.edu/~bates/JuliaForRProgrammers.pdf. The task is to use Gibbs sampler to sample from bivariate density

$$f(x,y) = kx^{2} \exp(-xy^{2} - y^{2} + 2y - 4x), x > 0,$$

using the conditional distributions

$$X|Y \sim \Gamma\left(3, \frac{1}{y^2 + 4}\right)$$

 $Y|X \sim \mathcal{N}\left(\frac{1}{1+x}, \frac{1}{2(1+x)}\right).$

Let's sample 10,000 points from this distribution with a thinning of 500.

- How long does R take? 42 seconds. http://hua-zhou.github.io/teaching/biostatm280-2016winter/gibbs_r.html
- How long does Julia take? 0.43 seconds. http://hua-zhou.github.io/teaching/biostatm280-2016winter/gibbs_julia.html
- With similar coding efforts, Julia offers ~ 100 fold speed-up! JIT in R didn't kick in. Neither does Matlab, which took about 20 seconds.
- Julia offers the capability of strong typing of variables. This facilitates the optimization by compiler.
- With little extra efforts, we can do parallel and distributed computing using Julia.

Benchmark of the same example in other languages including Rcpp is available in the blogs by Darren Wilkinson (http://bit.ly/IWhJ52) and Dirk Eddelbuettel's (http://dirk.eddelbuettel.com/blog/2011/07/14/).

"As some of you may know, I have had a (rather late) mid-life crisis and run off with another language called Julia. http://julialang.org"

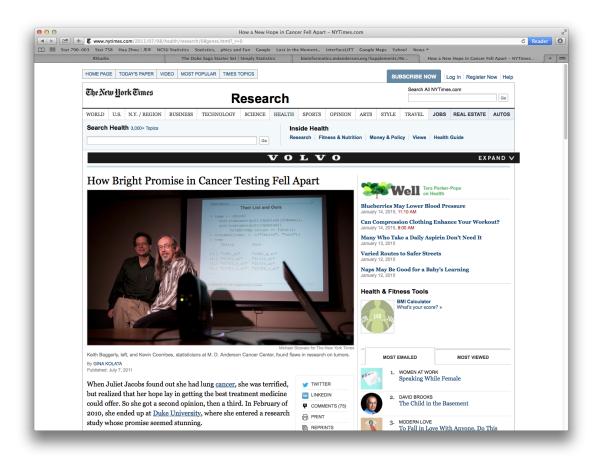
Doug Bates (on the knitr Google Group)

Reproducible research (in computational science)

An article about computational result is **advertising**, **not** scholarship. The actual scholarship is the full software environment, code and data, that produced the result.

Buckheit and Donoho (1995) also see Claerbout and Karrenbach (1992)

- 3 stories of *not* being reproducible.
 - Duke Potti Scandal.



Potti et al. (2006) Genomic signatures to guide the use of chemotherapeutics, *Nature Medicine*, 12(11):1294–1300.

Baggerly and Coombes (2009) Deriving chemosensitivity from cell lines: Forensic bioinformatics and reproducible research in high-throughput biology, *Ann. Appl. Stat.*, 3(4):1309–1334. http://projecteuclid.org/euclid.aoas/1267453942

More information is available at

http://en.wikipedia.org/wiki/Anil_Potti

http://simplystatistics.org/2012/02/27/the-duke-saga-starter-set/

- Nature Genetics (2013 Impact Factor: 29.648). 20 articles about microarray profiling published in *Nature Genetics* between Jan 2005 and Dec 2006.

Repeatability of published microarray gene expression analyses

John P A Ioannidis $^{1-3}$, David B Allison 4 , Catherine A Ball 5 , Issa Coulibaly 4 , Xiangqin Cui 4 , Aedín C Culhane 6,7 , Mario Falchi 8,9 , Cesare Furlanello 10 , Laurence Game 11 , Giuseppe Jurman 10 , Jon Mangion 11 , Tapan Mehta 4 , Michael Nitzberg 5 , Grier P Page 4,12 , Enrico Petretto 11,13 & Vera van Noort 14

Given the complexity of microarray-based gene expression studies, guidelines encourage transparent design and public data availability. Several journals require public data deposition and several public databases exist. However, not all data are publicly available, and even when available, it is unknown whether the published results are reproducible by independent scientists. Here we evaluated the replication of data analyses in 18 articles on microarray-based gene expression profiling published in *Nature Genetics* in 2005–2006. One table or figure from each article was independently evaluated by two teams of analysts. We reproduced two analyses in principle

research, the Uniform Guidelines of the International Committee of Medical Journal Editors state that authors should "identify the methods, apparatus and procedures in sufficient detail to allow other workers to reproduce the results". Making primary data publicly available has many challenges but also many benefits 13, Public data availability allows other investigators to confirm the results of the original authors, exactly replicate these results in other studies and try alternative analyses to see whether results are robust and to learn new things. Journals such as Nature Genetics require public data deposition as a prerequisite for publication for microarray-based research. Yet, the extent to which data are indeed made fully and accurately publicly available and permit con-

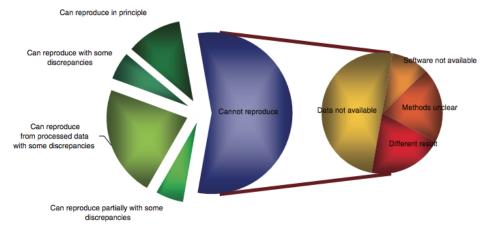
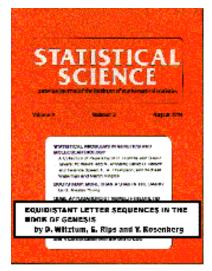
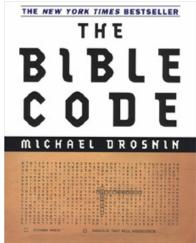


Figure 1 Summary of the efforts to replicate the published analyses.

- Bible code.





Witztum et al. (1994) Equidistant letter sequences in the book of genesis. *Statist. Sci.*, 9(3):429438. http://projecteuclid.org/euclid.ss/1177010393

McKay et al. (1999) Solving the Bible code puzzle, *Statist. Sci.*, 14(2):150–173.

https://www.math.washington.edu/~greenber/BibleCode.html

• Why reproducible research?

- Replicability has been a foundation of science. It helps accumulate scientific knowledge.
- Better work habit boosts quality of research.
- Greater research impact.
- Better teamwork. For you, it means better communication with your advisor (Buckheit and Donoho, 1995).

— ...

• Readings.

 Buckheit and Donoho (1995) Wavelab and reproducible research, in Wavelets and Statistics, volume 103 of Lecture Notes in Statistics, page 55–81. Springer Newt York. http://statweb.stanford.edu/~donoho/Reports/1995/wavelab.pdf

Donoho (2010) An invitation to reproducible computational research, *Biostatistics*, 11(3):385-388.

Peng (2009) Reproducible research and biostatistics, Biostatistics, 10(3):405–408.

Peng (2011) Reproducible research in computational science, *Science*, 334(6060):1226–1227.

Roger Peng's blogs Treading a New Path for Reproducible Research.

http://simplystatistics.org/2013/08/21/treading-a-new-path-for-reproducible-http://simplystatistics.org/2013/08/28/evidence-based-data-analysis-treadinghttp://simplystatistics.org/2013/09/05/implementing-evidence-based-data-analysis-treadinghttp://simplystatistics.org/2013/09/05/implementing-evidence-based-data-analysis-treadinghttp://simplystatistics.org/2013/09/05/implementing-evidence-based-data-analysis-treadinghttp://simplystatistics.org/2013/09/05/implementing-evidence-based-data-analysis-treadinghttp://simplystatistics.org/2013/09/05/implementing-evidence-based-data-analysis-treadinghttp://simplystatistics.org/2013/09/05/implementing-evidence-based-data-analysis-treadinghttp://simplystatistics.org/2013/09/05/implementing-evidence-based-data-analysis-treadinghttp://simplystatistics.org/2013/09/05/implementing-evidence-based-data-analysis-treadinghttp://simplystatistics.org/2013/09/05/implementing-evidence-based-data-analysis-treadinghttp://simplystatistics.org/2013/09/05/implementing-evidence-based-data-analysis-treadinghttp://simplystatistics.org/2013/09/05/implementing-evidence-based-data-analysis-treadinghttp://simplystatistics.org/2013/09/05/implementing-evidence-based-data-analysis-treadinghttp://simplystatistics.org/2013/09/05/implementing-evidence-based-data-analysis-treadinghttp://simplystatistics.org/2013/09/05/implementing-evidence-based-data-analysis-treadinghttp://simplystatis-treadinghttp

- Reproducible research with R and RStudio by Christopher Gandrud. It covers many useful tools: R, RStudio, LaTeX, Markdown, knitr, Github, Linux shell, ...

This book is nicely reproducible. Git clone the source from https://github.com/christophergandrud/Rep-Res-Book and you should be able to compile into a pdf.

- Reproducibility in Science at http://ropensci.github.io/reproducibility-guide/
- How to be reproducible in statistics?

When we publish articles containing figures which were generated by computer, we also publish the complete software environment which generates the figures.

Buckheit and Donoho (1995)

- For theoretical results, include all detailed proofs.
- For data analysis or simulation study
 - * Describe your computational results with painstaking details.

- * Put your code on your website or in an online supplement (required by many journals, e.g., *Biostatistics*, *JCGS*, ...) that allow replication of entire analysis or simulation study. A good example:
 - http://stanford.edu/~boyd/papers/admm_distr_stats.html
- * Create a dynamic version of your simulation study/data analysis.
- What can we do now? At least make your homework reproducible!
 - Document everything!
 - Everything is a text file (.csv, .tex, .bib, .Rmd, .R, ...) They aid future proof and are subject to version control.
 - Word/Excel are not text files.
 - All files should be human readable. Abundant comments and adopt a good style.
 - Tie your files together.
 - Use a dynamic document generation tool (weaving/knitting text, code, and output together) for documentation.
 - Use a version control system proactively.
 - Print sessionInfo() in R.

For your homework, submit (put in the master branch) a final pdf or html report and all files and instructions necessary to reproduce all results.

- Tools for dynamic document/report generation.
 - R: RMarkdown, knitr, Sweave.
 - Matlab: automatic report generator.
 - Python: IPython, Pweave.
 - Julia: IJulia.

For this course, please write homework report in RMarkdown (or IJulia if you use Julia).

4 Lecture 4, Jan 14

Last time

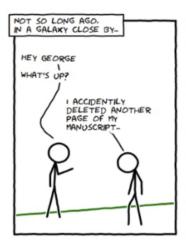
- Computer languages.
- Comparison of R, MATLAB, and JULIA.
- R, RStudio, RMarkdown.

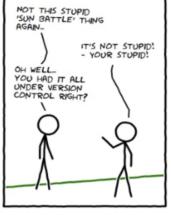
Today

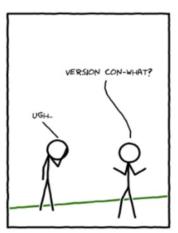
• Version control.

Version control by Git

If it's not in source control, it doesn't exist.







- Collaborative research. Statisticians, as opposed to "closet mathematicians", rarely do things in vacuum.
 - We talk to scientists/clients about their data and questions.
 - We write code (a lot!) together with team members or coauthors.
 - We run code/program on different platforms.
 - We write manuscripts/reports with co-authors.

– ...

- 4 things distinguish professional programmers from amateurs:
 - Use a version control system.
 - Automate repetitive taks.
 - Systematic testing.
 - Use debugging aids rather than print statements.

• Why version control?

- A centralized repository helps coordinate multi-person projects.
- Time machine. Keep track of all the changes and revert back easily (reproducible).
- Storage efficiency.
- Synchronize files across multiple computers and platforms.
- github.com is becoming a de facto central repository for open source development.
 E.g., all packages in Julia are distributed through github.com.
- Advertise yourself thru github.com.
- Available version control tools.
 - Open source: cvs, subversion (aka svn), Git, ...
 - Proprietary: Visual SourceSafe (VSS), ...
 - Dropbox? Mostly for file back and sharing, limited version control (1 month?), ...

We use Git in this course.

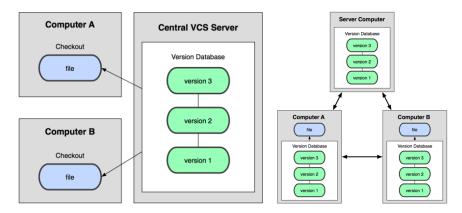
• Why Git?

- The Eclipse Community Survey in 2014 shows Git is the most widely used source code management tool now. Git (33.3%) vs svn (30.7%).
- History: Initially designed and developed by Linus Torvalds in 2005 for Linux kernel development. "git" is the British English slang for "unpleasant person".

I'm an egotistical bastard, and I name all my projects after myself. First 'Linux', now 'git'.

Linus Torvalds

 A fundamental difference between svn (centralized version control system, left plot) and Git (distributed version control system, right plot):



- Advantages of Git.
 - * Free and open source.
 - * Speed and simple (?) design.
 - * Strong support for non-linear development (1000s of parallel branches). Scalable to large projects like the Linux kernel project.
 - * Fully distributed. Fast, no internet required, disaster recovery,
- Be aware that svn is still widely used in IT industry (Apache, GCC, Source-Forge, Google Code, ...) and R development. E.g., type
 svn log -v -l 5 https://svn.r-project.org/R
 on command line to get a glimpse of what R development core team is doing. Good to master some basic svn commands.
- What do I need to use Git?
 - A Git server enabling multi-person collaboration through a centralized repository.
 - * github.com: unlimited public repositories, private repositories costs \$, academic user can get 5 private repositories for free.

- * bitbucket.org: unlimited private repositories for academic account (register for free using your UCLA email).
- * Some institutes and departments have their own git server.

We use bitbucket.org in this course.

- Git client.
 - * Linux: installed on many servers. If not, install on CentOS by yum install git.
 - * Mac: install by port install git.
 - * Windows: GitHub for Windows (GUI), TortoiseGIT (is this good?)

Don't rely on GUI. Learn to use Git on command line.

• Life cycle of a (research) project.

Stage 1:

- A project (idea) is started on bitbucket.org, with directories say codebase,
 datasets, manuscripts, talks, ...
- Advantage of bitbucket.org: privacy of research ideas (free private repositories).
- Downside of bitbucket.org: not as widely known as github.com.

Stage 2:

- Hopefully, research idea pans out and we want to distribute a standalone software development, e.g., an R package, repository at github.com. Read the (free) book R Packages (http://r-pkgs.had.co.nz) by Hadley Wickham for development of R packages and distribution via github.com.
- This usually inherits from the codebase folder and happens when we submit a paper.
- Challenges: keep all version history. It's doable.

Stage 3:

- Active maintenance of the public software repository.

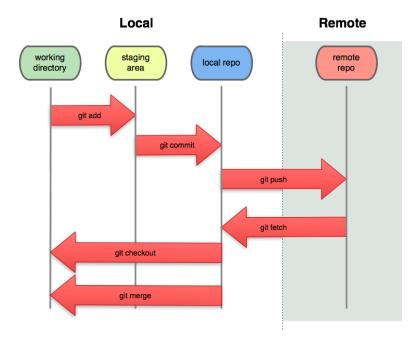
- At least three branches: develop, master, gh-pages.

develop: main development area.

master: software release.

gh-pages: software webpage.

• Basic workflow of Git.



- Synchronize local Git directory with remote repository (git pull).
- Modify files in local working directory.
- Add snapshots of them to staging area (git add).
- Commit: store snapshots permanently to (local) Git repository (git commit).
- Push commits to remote repository (git push).

• Basic Git usage.

- Register for an account on a Git server, e.g., bitbucket.org. Fill out your profile, upload your public key to the server, ...
- Identify yourself at local machine:
 git config --global user.name "Hua Zhou"
 git config --global user.email "huazhou@ucla.edu"
 Name and email appear in each commit you make.

- Initialize a project:
 - * Create a repository, e.g., biostat-m280-2016-winter, on the server bitbucket.org. Then clone to local machine git clone git@bitbucket.org:username/biostat-m280-2016-winter.git
 - * Alternatively use following commands to initialize a Git directory from a local folder and then push to the Git server git init

git remote add origin git@bitbucket.org:username/biostat-m280-2016-winter git push -u origin master

- Edit working directory.

git pull update local Git repository with remote repository (fetch + merge).

git status displays the current status of working directory.

git log filename displays commit logs of a file.

git diff shows differences (by default difference from the most recent commit).

git add ... adds file(s) to the staging area.

git commit commits changes in staging area to Git directory.

git push publishes commits in local Git directory to remote repository.

Following demo session is on my local Mac machine.

```
hzhou3@Hua-Zhous-MacBook-Pro:mglm $ pwd
/Users/hzhou3/github.ncsu/mglm
hzhou3@Hua-Zhous-MacBook-Pro:mglm $ ls
.DS_Store .gitignore
                        datasets/
                                      manuscripts/
                         literature/ talks/
            codebase/
.ait/
hzhou3@Hua-Zhous-MacBook-Pro:mglm $ git pull
remote: Counting objects: 5, done.
remote: Compressing objects: 100% (2/2), done.
remote: Total 5 (delta 3), reused 5 (delta 3)
Unpacking objects: 100% (5/5), done.
From github.ncsu.edu:hzhou3/vctest
   80be212..b22d29f master
                               -> origin/master
Updating 80be212..b22d29f
Fast-forward
 manuscripts/letter-skat-famskat/Letter_to_the_editor.tex | 4 ++--
 1 file changed, 2 insertions(+), 2 deletions(-)
hzhou3@Hua-Zhous-MacBook-Pro:mglm $ echo "hello st790 class" > gitdemo txt
hzhou3@Hua-Zhous-MacBook-Pro:mglm $ ls
.DS_Store .gitignore datasets/
                                      literature/ talks/
                         gitdemo.txt manuscripts/
.git/
            codebase/
```

```
hzhou3@Hua-Zhous-MacBook-Pro:mglm $ git add gitdemo.txt
 hzhou3@Hua-Zhous-MacBook-Pro:mglm $ git status .
 On branch master
 Your branch is up-to-date with 'origin/master'.
 Changes to be committed:
    (use "git reset HEAD <file>..." to unstage)
    new file: gitdemo.txt
 Untracked files:
    (use "git add <file>..." to include in what will be committed)
    codebase/Example_RNAseq_top100/
    codebase/MGLM/R/.Rhistory
 hzhou3@Hua-Zhous-MacBook-Pro:mglm $ git commit -m "git demo for st790 class"
  [master ea636ff] git demo for st790 class
  1 file changed, 1 insertion(+)
  create mode 100644 gitdemo.txt
 hzhou3@Hua-Zhous-MacBook-Pro:mglm $ git log gitdemo.txt
 commit ea636ff5665bc26bf8a79751b75d0e9d67bdb7d1
 Author: Hua Zhou <hua_zhou@ncsu.edu>
         Sun Jan 11 17:06:23 2015 -0500
     git demo for st790 class
 hzhou3@Hua-Zhous-MacBook-Pro:mglm $ git push
 Counting objects: 3, done.
 Delta compression using up to 8 threads.
 Compressing objects: 100% (2/2), done.
 Writing objects: 100% (3/3), 301 bytes | 0 bytes/s, done. Total 3 (delta 1), reused 0 (delta 0)
 To git@github.ncsu.edu:hzhou3/mglm.git
     77145d2..ea636ff master -> master
git reset --soft HEAD 1 undo the last commit.
git checkout filename go back to the last commit.
git rm different from rm.
Although git rm deletes files from working directory. They are still in Git
history and can be retrieved whenever needed. So always be cautious to
put large data files or binary files into version control.
```

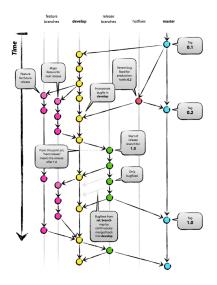
```
hzhou3@Hua-Zhous-MacBook-Pro:mglm $ echo "bye st790 class" >> gitdemo.txt
hzhou3@Hua-Zhous-MacBook-Pro:mglm $ cat gitdemo.txt
hello st790 class
bye st790 class
hzhou3@Hua-Zhous-MacBook-Pro:mglm $ git diff gitdemo.txt
diff --git a/gitdemo.txt b/gitdemo.txt
index ece6d4e..2bb77f8 100644
--- a/gitdemo.txt
+++ b/gitdemo.txt
 hello st790 class
+bve st790 class
hzhou3@Hua-Zhous-MacBook-Pro:mglm $ git checkout gitdemo.txt
hzhou3@Hua-Zhous-MacBook-Pro:mglm $ cat gitdemo.txt
hello st790 class
hzhou3@Hua-Zhous-MacBook-Pro:mglm $ git rm gitdemo.txt
rm 'gitdemo.txt'
hzhou3@Hua-Zhous-MacBook-Pro:mglm $ git status .
On branch master
Your branch is up-to-date with 'origin/master'.
Changes to be committed:
  (use "git reset HEAD <file>..." to unstage)
   deleted:
                 gitdemo.txt
Untracked files:
  (use "git add <file>..." to include in what will be committed)
   codebase/Example_RNAseq_top100/
   codebase/MGLM/R/.Rhistory
hzhou3@Hua-Zhous-MacBook-Pro:mglm $ git commit -m "delete the git demo file for st790"
[master 4b4f9c5] delete the git demo file for st790
 1 file changed, 1 deletion(-)
 delete mode 100644 gitdemo.txt
hzhou3@Hua-Zhous-MacBook-Pro:mglm $ git push
Counting objects: 2, done.
Delta compression using up to 8 threads.

Compressing objects: 100% (2/2), done.

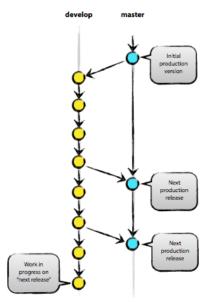
Writing objects: 100% (2/2), 238 bytes | 0 bytes/s, done.

Total 2 (delta 1), reused 0 (delta 0)
lTo git@github.ncsu.edu:hzhou3/mglm.git
ea636ff..4b4f9c5 master -> master
hzhou3@Hua-Zhous-MacBook-Pro:mglm $ ls
.DS_Store .gitignore datasets/
                                              manuscripts/
.git/
                               literature/ talks/
               codebase/
```

- Branching in Git.
 - Branches in a project:



- For this course, you need to have two branches: develop for your own development and master for releases (homework submission). Note master is the default branch when you initialize the project; create and switch to develop branch immediately after project initialization.



- Commonly used commands:
 - $\mbox{git branch branch name} \ \mbox{creates a branch}.$
 - git branch shows all project branches.
 - git checkout branchname switches to a branch.
 - git tag shows tags (major landmarks).

git tag tagname creates a tag.

- Let's look at a typical branching and merging workflow.
 - * Now there is a bug in v0.0.3 \dots

```
hzhou3@Hua-Zhous-MacBook-Pro:gitdemo $ git branch develop

* master
hzhou3@Hua-Zhous-MacBook-Pro:gitdemo $ git tag
v0.0.1
v0.0.2
v0.0.3
hzhou3@Hua-Zhous-MacBook-Pro:gitdemo $ ls
.git/ bug.txt code.txt
hzhou3@Hua-Zhous-MacBook-Pro:gitdemo $
```

How to organize version number of your software? Read blog "R Package Versioning" by Yihui Xie

http://yihui.name/en/2013/06/r-package-versioning/

```
000
                                gitdemo - bash - 80×31
hzhou3@Hua-Zhous-MacBook-Pro:gitdemo $ git checkout develop
Switched to branch 'develop'
Your branch is up-to-date with 'origin/develop'.
hzhou3@Hua-Zhous-MacBook-Pro:gitdemo $ git branch
* develop
 master
hzhou3@Hua-Zhous-MacBook-Pro:gitdemo $ ls
.git/
         code.txt
hzhou3@Hua-Zhous-MacBook-Pro:gitdemo $ git pull origin master
From github.ncsu.edu:hzhou3/gitdemo
                     master
                                -> FETCH_HEAD
* branch
Updating 44dd1d1..da047cf
Fast-forward
bug.txt | 1 +
1 file changed, 1 insertion(+)
create mode 100644 bug.txt
hzhou3@Hua-Zhous-MacBook-Pro:gitdemo $ ls
         bug.txt code.txt
.git/
hzhou3@Hua-Zhous-MacBook-Pro:gitdemo $ git rm bug.txt
rm 'bug.txt'
hzhou3@Hua-Zhous-MacBook-Pro:gitdemo $ git commit -m "debug"
[develop 1085b4c] debug
1 file changed, 1 deletion(-)
delete mode 100644 bug.txt
hzhou3@Hua-Zhous-MacBook-Pro:gitdemo $ git push
Counting objects: 1, done.
Writing objects: 100% (1/1), 180 bytes | 0 bytes/s, done.
Total 1 (delta 0), reused 0 (delta 0)
To git@github.ncsu.edu:hzhou3/gitdemo.git
   44dd1d1..1085b4c develop -> develop
hzhou3@Hua-Zhous-MacBook-Pro:gitdemo $
```

Now 'debug' in develop branch is ahead of master branch.

* Merge bug fix to the master branch.

```
000
                               gitdemo — bash — 80×26
hzhou3@Hua-Zhous-MacBook-Pro:gitdemo $ git checkout master
Switched to branch 'master'
Your branch is up-to-date with 'origin/master'.
hzhou3@Hua-Zhous-MacBook-Pro:gitdemo $ git branch
* master
hzhou3@Hua-Zhous-MacBook-Pro:gitdemo $ git pull origin develop
From github.ncsu.edu:hzhou3/gitdemo
* branch
                    develop -> FETCH_HEAD
Updating da047cf..1085b4c
Fast-forward
bug.txt | 1 -
1 file changed, 1 deletion(-)
delete mode 100644 bug.txt
hzhou3@Hua-Zhous-MacBook-Pro:gitdemo $ ls
        code.txt
.git/
hzhou3@Hua-Zhous-MacBook-Pro:gitdemo $ git status .
On branch master
Your branch is ahead of 'origin/master' by 1 commit.
  (use "git push" to publish your local commits)
nothing to commit, working directory clean
hzhou3@Hua-Zhous-MacBook-Pro:gitdemo $ git push origin master
Total 0 (delta 0), reused 0 (delta 0)
To git@github.ncsu.edu:hzhou3/gitdemo.git
   da047cf..1085b4c master -> master
hzhou3@Hua-Zhous-MacBook-Pro:gitdemo $
```

* Tag a new release v0.0.4.

```
gitdemo — bash — 80×26
000
hzhou3@Hua-Zhous-MacBook-Pro:gitdemo $ git tag v0.0.4
hzhou3@Hua-Zhous-MacBook-Pro:gitdemo $ git tag
v0.0.1
v0.0.2
v0.0.3
v0.0.4
hzhou3@Hua-Zhous-MacBook-Pro:gitdemo $ git show v0.0.4
commit 1085b4c97ed29fc847442bd0640db2b6fed4d0af
Author: Hua Zhou <hua_zhou@ncsu.edu>
Date:
       Tue Jan 13 11:24:19 2015 -0500
    debug
diff --git a/bug.txt b/bug.txt
deleted file mode 100644
index 0ald6ac..0000000
 -- a/bug.txt
+++ /dev/null
@ -1 +0,0 @
-There is a bug
hzhou3@Hua-Zhous-MacBook-Pro:gitdemo $ git push origin v0.0.4
Total 0 (delta 0), reused 0 (delta 0)
To git@github.ncsu.edu:hzhou3/gitdemo.git
* [new tag]
                    v0.0.4 -> v0.0.4
hzhou3@Hua-Zhous-MacBook-Pro:gitdemo $
```

• Further resources:

- Book Pro Git, http://git-scm.com/book/en/v2
- Google
- Some etiquettes of using Git and version control systems in general.
 - Be judicious what to put in repository
 - * Not too less: Make sure collaborators or yourself can reproduce everything on other machines.
 - * Not too much: No need to put all intermediate files in repository.

Strictly version control system is for source files only. E.g. only xxx.tex, xxx.bib, and figure files are necessary to produce a pdf file. Pdf file doesn't need to be version controlled or frequently committed.

- "Commit early, commit often and don't spare the horses"

Adding an informative message when you commit is not optional. Spending one minute now saves hours later for your collaborators and yourself.
 Read the following sentence to yourself 3 times:

"Write every commit message like the next person who reads it is an axewielding maniac who knows where you live."

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