Wednesday, January 24, 2018 12:44 PM

Call a method 's a first-order Hethod is each iteration "it uses all previous couputed quadrants  $\nabla F(X, 1), \nabla F(X, 5), \dots, \nabla F(X_{K-1})$ and initial X,  $F(x_{1}), F(x_{2}), \dots, F(x_{k-1})$ to generate a new point XK. In short, Frist-order Method can only access to 5 and DF. Remark: These are other acess nudel such as 5t and 75t. Usually, East-oslar method satisfies XK & Gpan(XI, Rf(KI), Rf(K2), ..., Vf(XK-1)) According to the definition, cutting plane inthoses are first-order. However, the common use of the term "First-order method" seters to methods that only use "sruple formal" A typical First-order method? XK41 = XK - 7 VS(XK) (grachent descent) Common betters: First-order method (there with single tomber) works well for large-scale Problems My belief: There should be something that is better. Practicality of Cutting Hone Methods. How '14 'It possible to say consiste the carter of gravity of 1 billions downers -> I am writing a package with runny time closer to timeor. Let see how long dees 'it take. -> You naver need to do so. Note that all cutting plane mithod we mentioned satisfies  $X_{k} = \text{Span}(X_{1}, \text{VS}(X_{1}), \text{VS}(X_{2}), \dots, \text{VS}(X_{k-1})) \quad (k \mid dm)$ Therefore, when we only need to solve some K-1 chim problem at Kth dep.

Therefore, whey, we only read to solve some K-1 chim problem at Kth step. Roughly spealing, is the cattery plane method involves solving a problem that takes F(k) time for IRK. Then, the total time for the 1th iter is T(VF) + NK + F(K) compute gradrait mention kolm solve some avery subspace problem As I mentioned last time, for pooblem of the form 25; (a; x-b; ), T(VF) is often dontrated by computery AV for some V where A= (a; So, a more bain runtime estimate for cutting plane method is nuter of nonzero Mnz(A) + nK + f(K) As conjourision, L-BifGs takes nuter of nonzero Mnz(A) + nK. It it is a large - scale problem,  $f(k) \ll u k$ . My belies: Cutting plane methods becomes more conjetitive as N > 020. Is the proton is lorge scale and we spent lots of time to get VS, Then it sounds silly to just combine these gradient by single combination. Note that IPM, CHOL, Muttigrid, AGD, they all takes decades to become recongized. Not sure about cutting plane methods. Thesis Question: Develope cathy the method that take adventuge of Z-5; structure and Implant it. Summy of First-order methods & stronly conver, B smith G-Lipsdiz, R= 1/xo-X"/2 Algorithm errors (Aune the function is Gradient Descent BR/4 2 équivalent BR2 (1- 3)t Hirror Descent GR/FE tight tight ) reduction Get Arrob-tal BR<sup>2</sup>/42 traht κ.,

TIGHT L Vit Accelerated Gradiest Decent tight 5 ", O(tight) ßR<sup>2</sup>/42 BR?(1-原)" tight  $(1-\frac{1}{n})^{R(t)}$ Certain Cuiting plane Methods be2(1-房)t O(tight) Misconceptron: linear convergence is better than subfram convergence (1-E)t We can always solve convergence (aka (1-m)-P(+)) Misconceptron: drunensson undependent result is touter for large scale problem. Even for very simple postdem like  $\sum (x_i - x_{i+1})^2$ , we have  $\frac{x}{2} = N^2$ . Theofore drumasin independent result can depends on drumesson. Relations between those bounds (1) Suppose we have a way to tool & st  $f(x) - f(x'') \in \frac{G'}{\alpha t}$ for any 02-strongly convex, Gi-Lip 5 For any Gi-tip convex F, we consider For (x)= F(x)+ & 1/x-xoll<sup>2</sup> We have that  $f(x) - f(x^{*}) \leq f_{\alpha}(x) - f_{\alpha}(x^{*}) + \stackrel{\sim}{\neq} ||x^{*} - y_{o}||^{2}$ < It + 2 R<sup>2</sup> (used the algo above) Chossing the best &, we have  $f(x) - f(x^*) \leq \frac{GP}{JZT}$ Hence, if we have an algorithm achieves Ge, we can them it to an algorithm achieves GR. Suppose we know BP- , IS F is a-straight convex, we have  $(\mathbf{z})$ 

Suppose we know BK , IS F is a straight convex, we have  $(\mathbf{Z})$  $\frac{2}{2}|x_{4}-x^{*}||^{2} \leq \frac{4}{2}(x_{4}) - \frac{4}{2}(x^{*}) \leq \frac{1}{2}|x_{0}-x^{*}||^{2}$ Set t= 2 €, we have 1/xy-x"112 ≤ ± 1/xo-x"112 Hence, every 2 B, the distance to optimum decrease by 7. By restarting the algorithm, we have BR2 (1-2 13) T Lower bounds Consider the protein: -X, + - - X, + - Z Z X? The soln is of the from ~c(1-层)' --> i they approximite solin has support at least I . Next, wit that is XK = Span(X, , VS(X,), VS(X,), ..., VF(Xe-1)) and 'IF X,= (1,0,0,...,0) Than XX has only & non-zeros in the beginning. Therefore, we need I iterations. All First-order methods including cutting plane methods has their serious issue # "ter > " drameter of the problem". Open problem: Is there any vary to help the connectivity for first How about

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