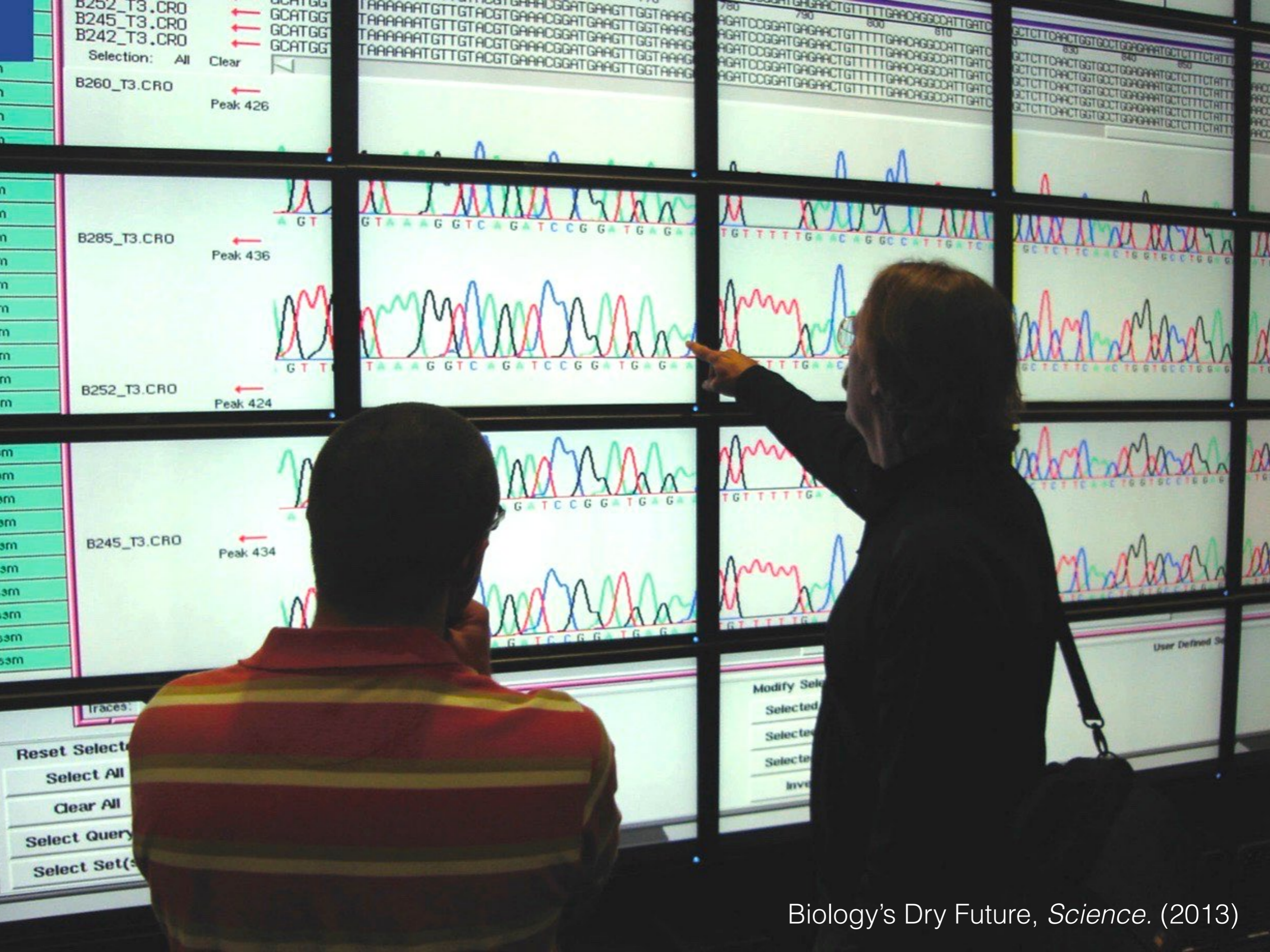


hdnom.io: High-Dimensional Survival Modeling with Shiny

Nan Xiao @road2stat

#ShinyDevCon



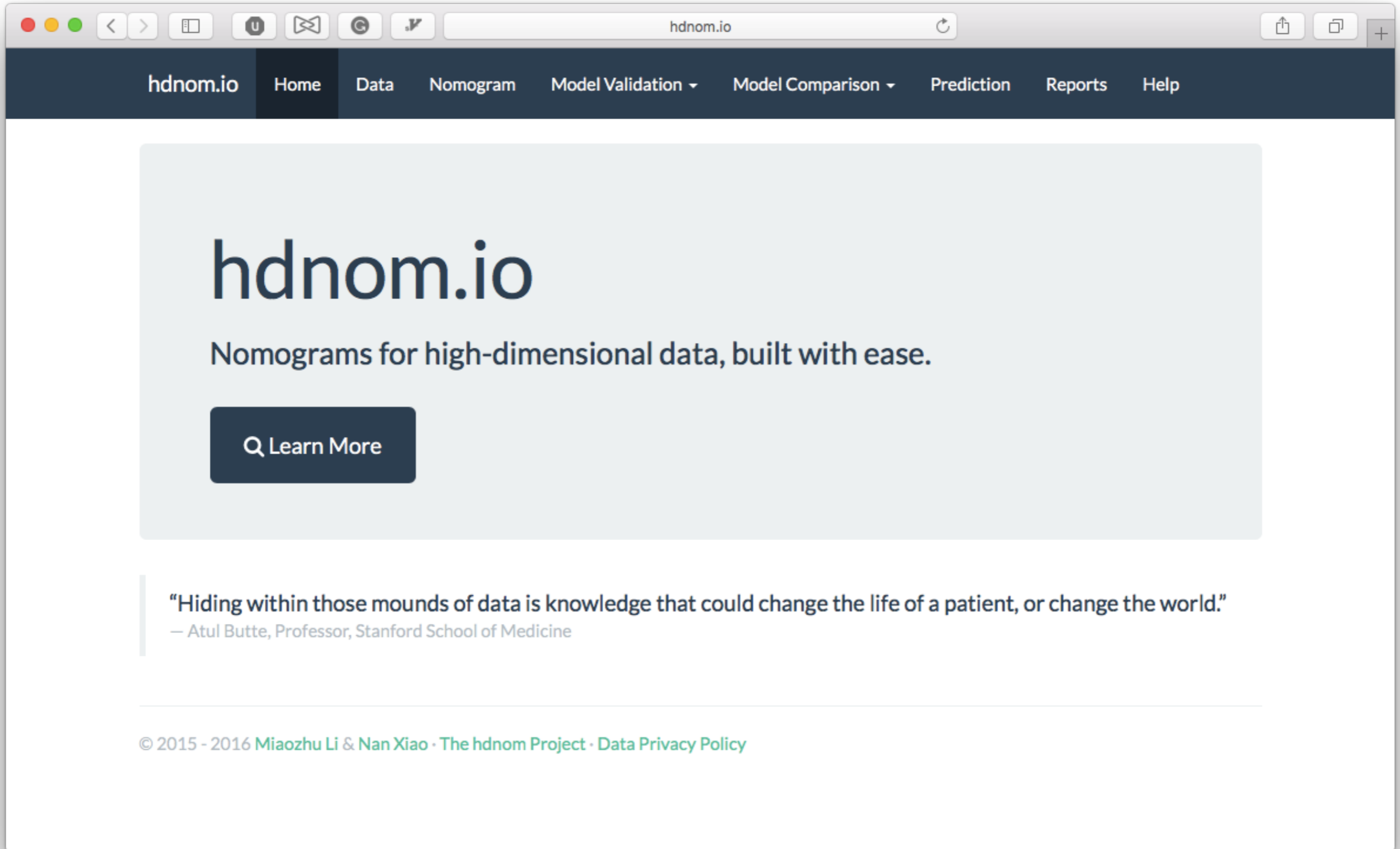
B252_T3.CRO
B245_T3.CRO
B242_T3.CRO
Selection: All
Clear
B260_T3.CRO
Peak 426

B285_T3.CRO
Peak 436
B252_T3.CRO
Peak 424

B245_T3.CRO
Peak 434

Traces:
Reset Select
Select All
Clear All
Select Query
Select Set(s)

Modify Sel
Selected
Selected
Selected
Inve



hdnom.io

Nomograms for high-dimensional data, built with ease.

🔍 Learn More

“Hiding within those mounds of data is knowledge that could change the life of a patient, or change the world.”

— Atul Butte, Professor, Stanford School of Medicine

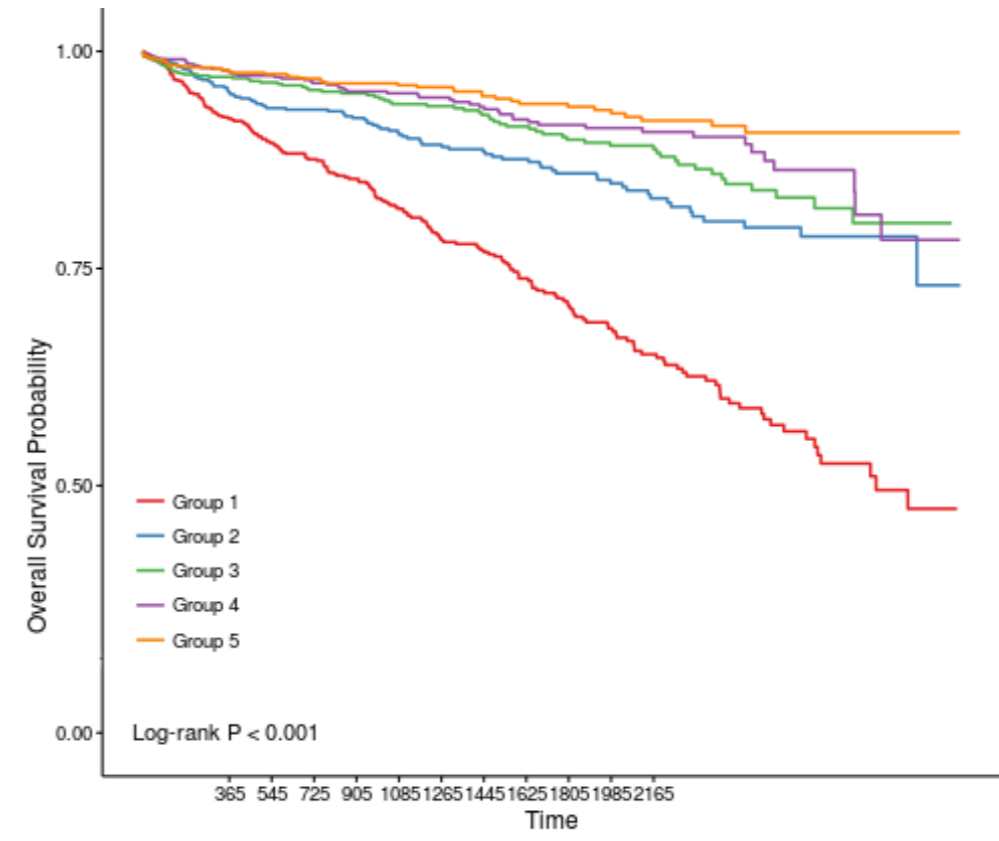
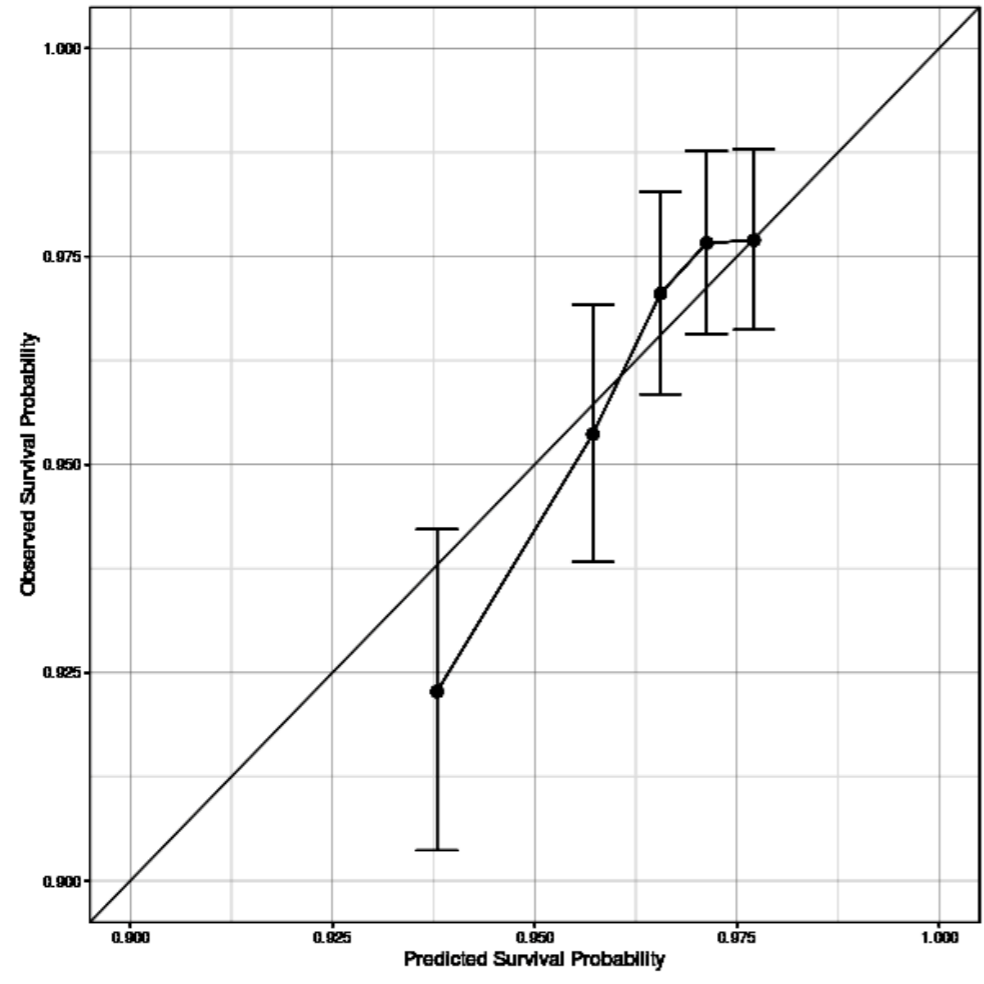
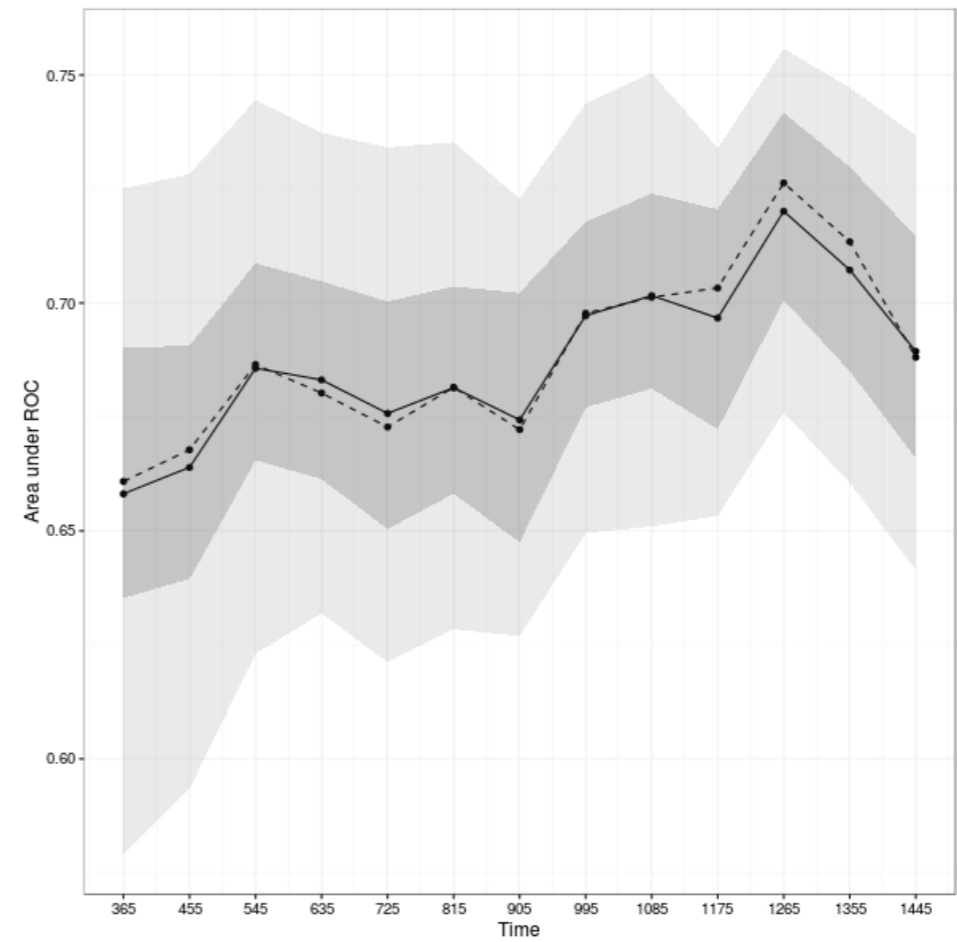
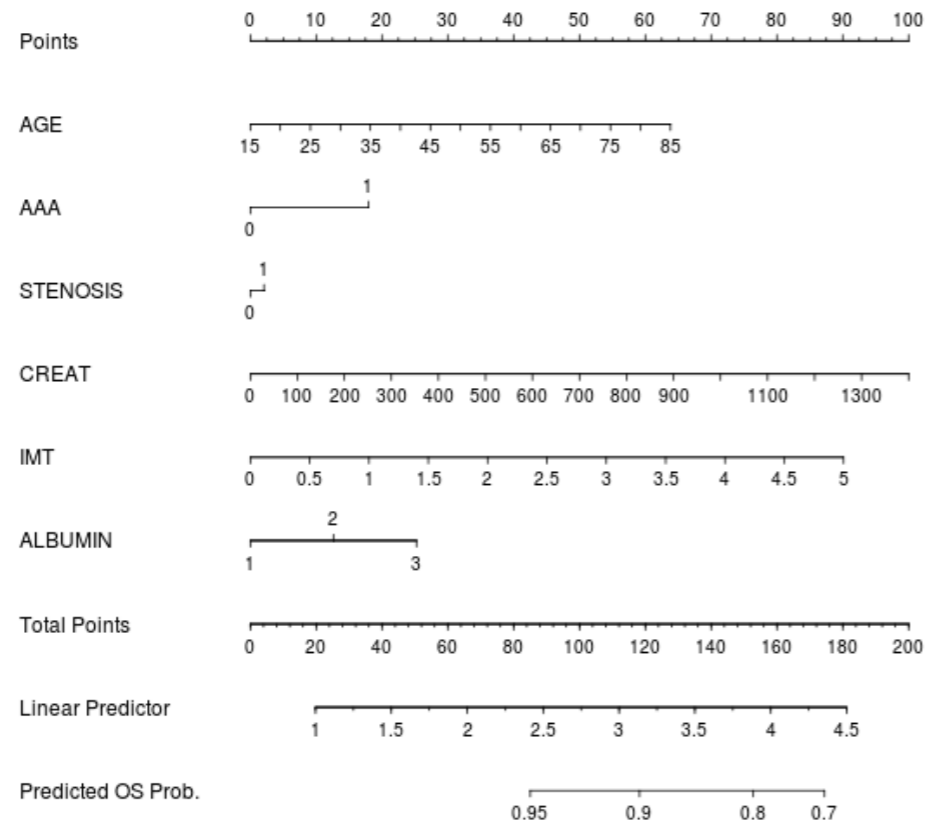
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Methods

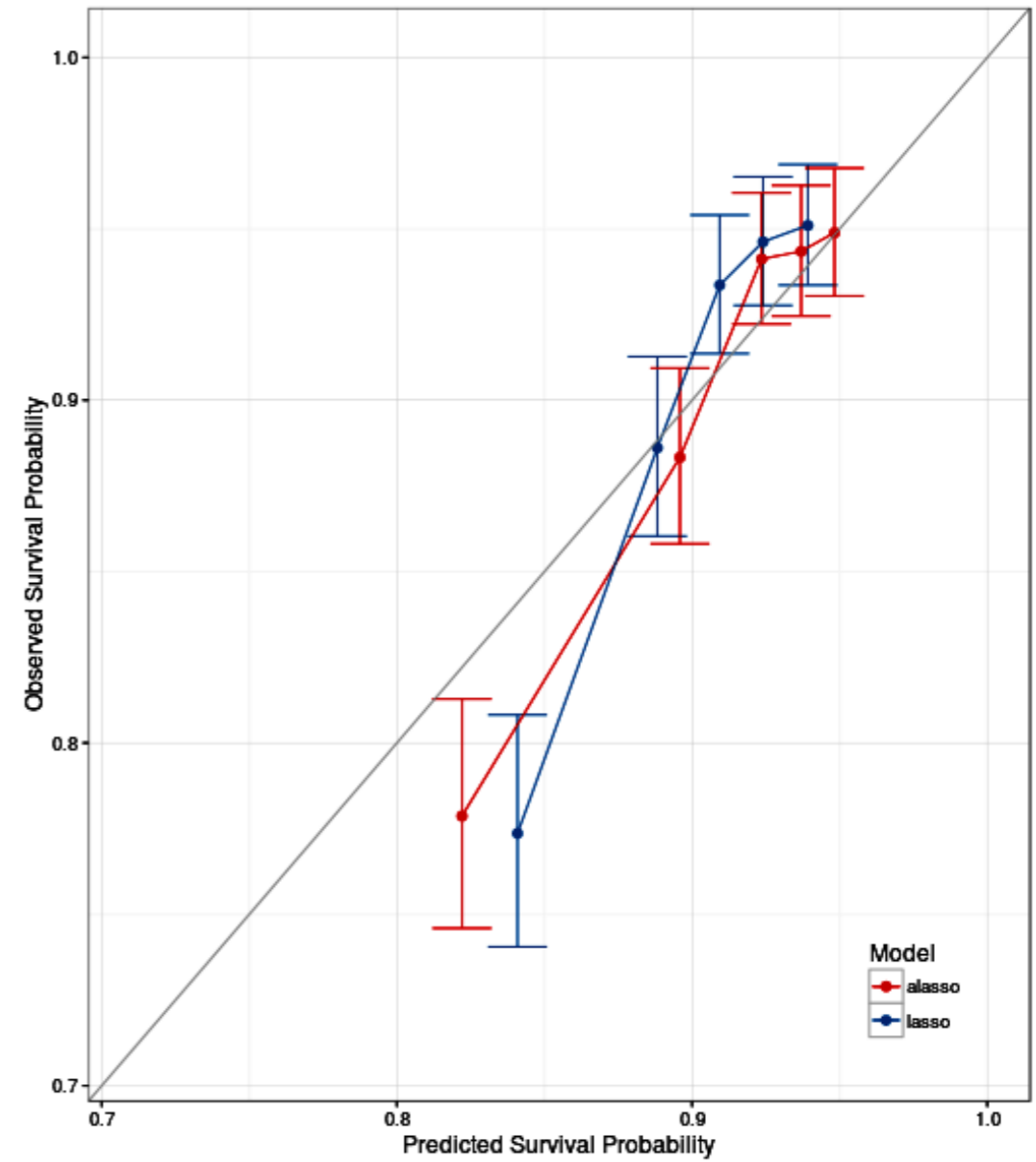
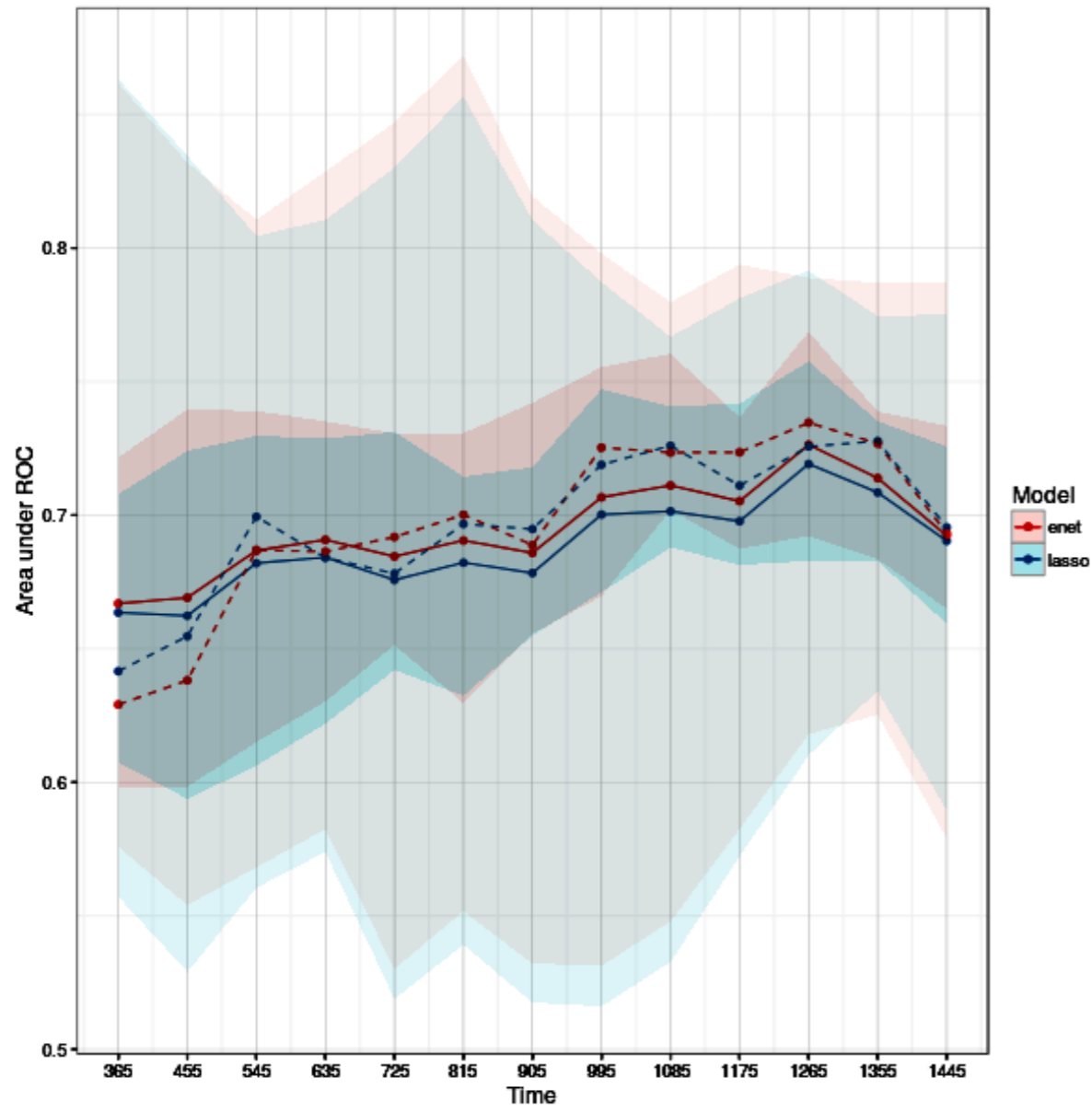
- Lasso / Adaptive Lasso / Fused Lasso
- Elastic-Net / Adaptive Elastic-Net
- SCAD / Snet
- MC+ / Mnet

Features

Basically all the survival modeling results you need to put in the research paper.



Group 5	637	585	540	485	443	394	348	308	274	237	212
Group 4	644	602	544	476	418	370	335	299	273	234	196
Group 3	637	580	523	478	440	398	359	314	282	251	218
Group 2	620	561	519	461	404	350	312	277	249	216	179
Group 1	638	593	525	471	406	354	323	275	232	192	164
	Number at risk										



Model Comparison

Demo

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Feedback:

miaozhu.li@duke.edu

nanx@uchicago.edu