



Week 12

Presentation

PHY 496

BRADEN KRONHEIM

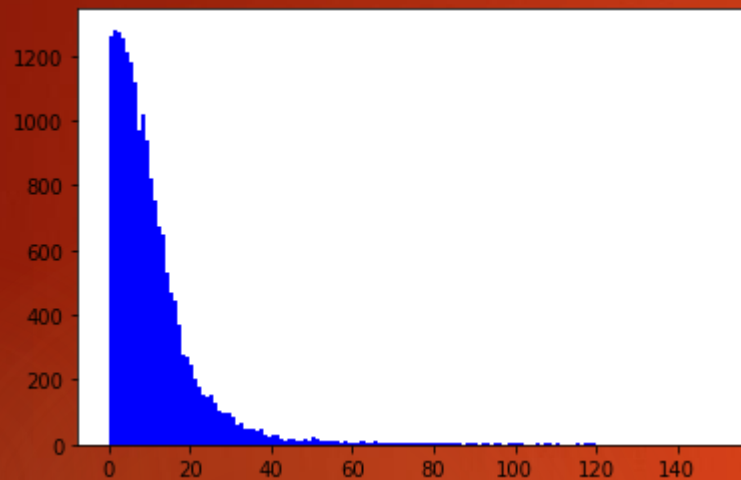
APRIL 12, 2019

Summary

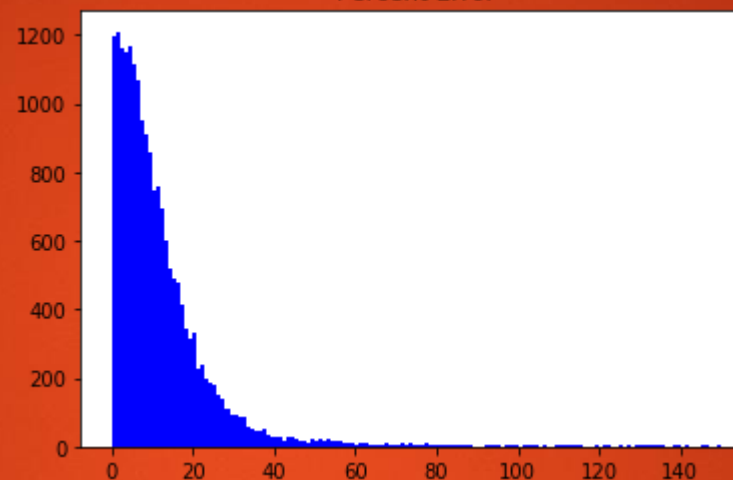
- ▶ Ran code to generate 50,000 networks, saving every 10
 - ▶ Code broke just before 40,000, still have 3,900 networks
- ▶ Started more general error analysis
 - ▶ Real vs. Predicted
 - ▶ Error from each input variable
- ▶ Started working on poster

General % Error

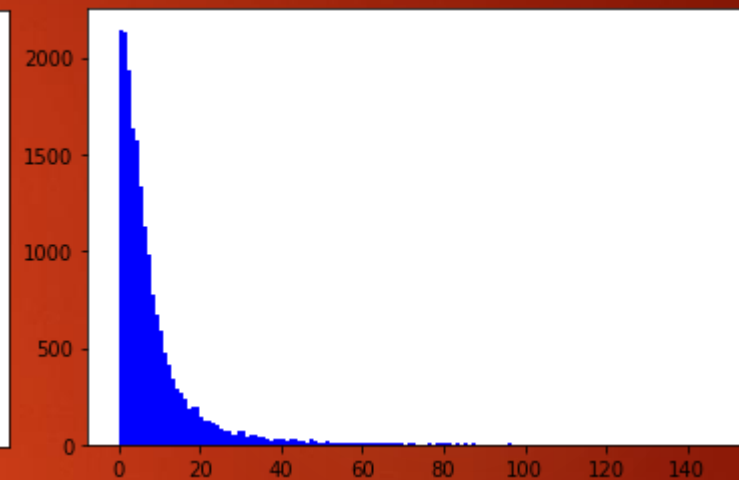
Flipout
Percent Error



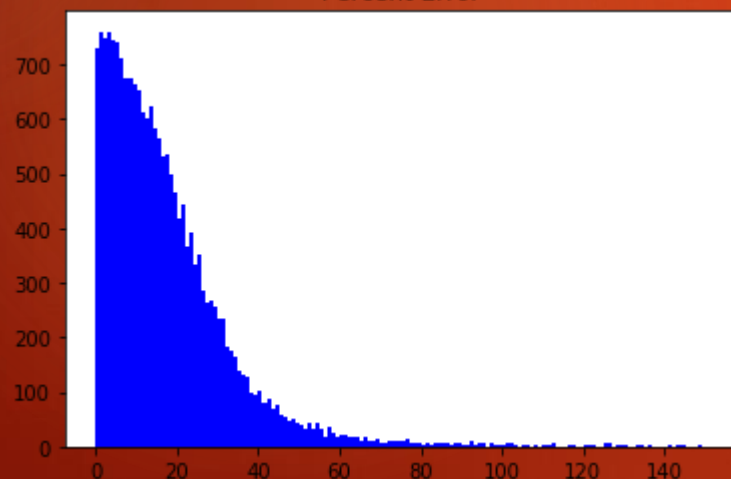
Reparameterization
Percent Error



HMC 5000
Percent Error

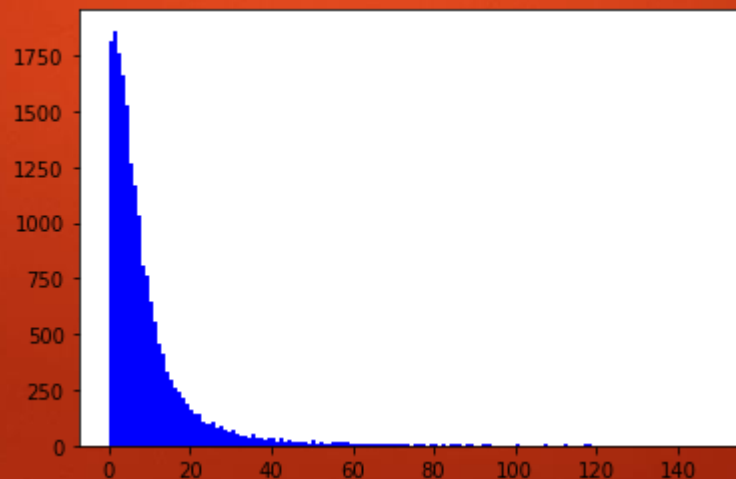


Normal
Percent Error



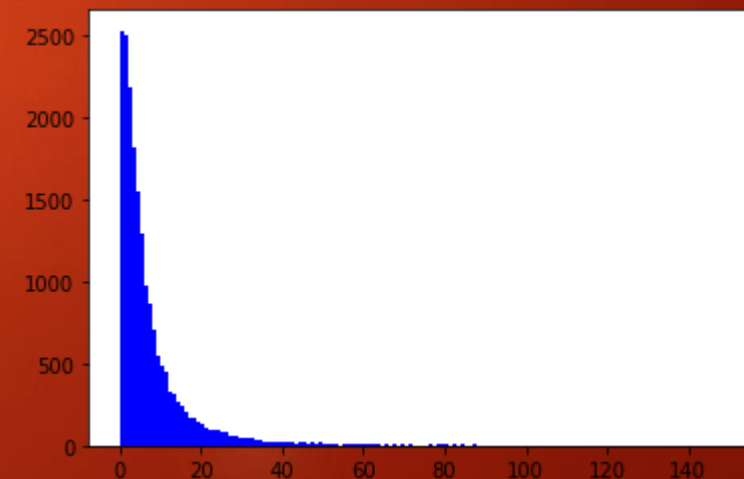
HMC 1000

Percent Error



HMC 195 from 3900

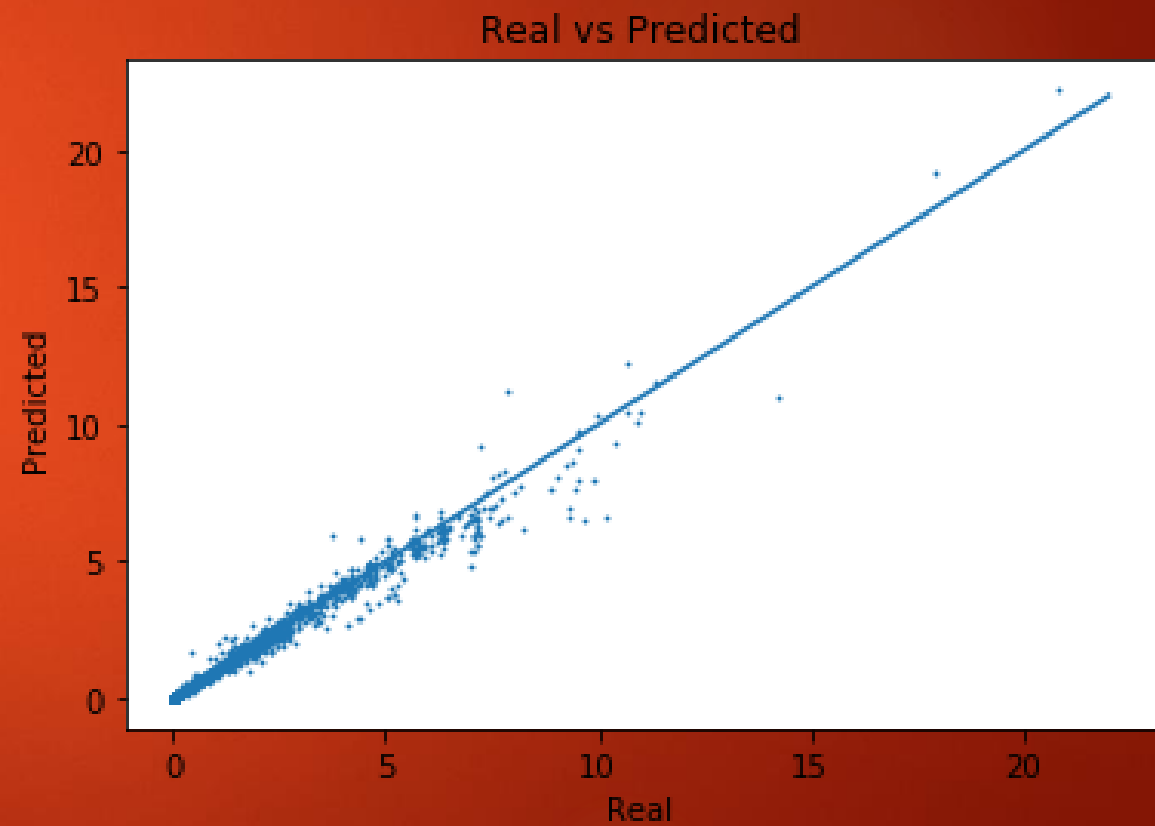
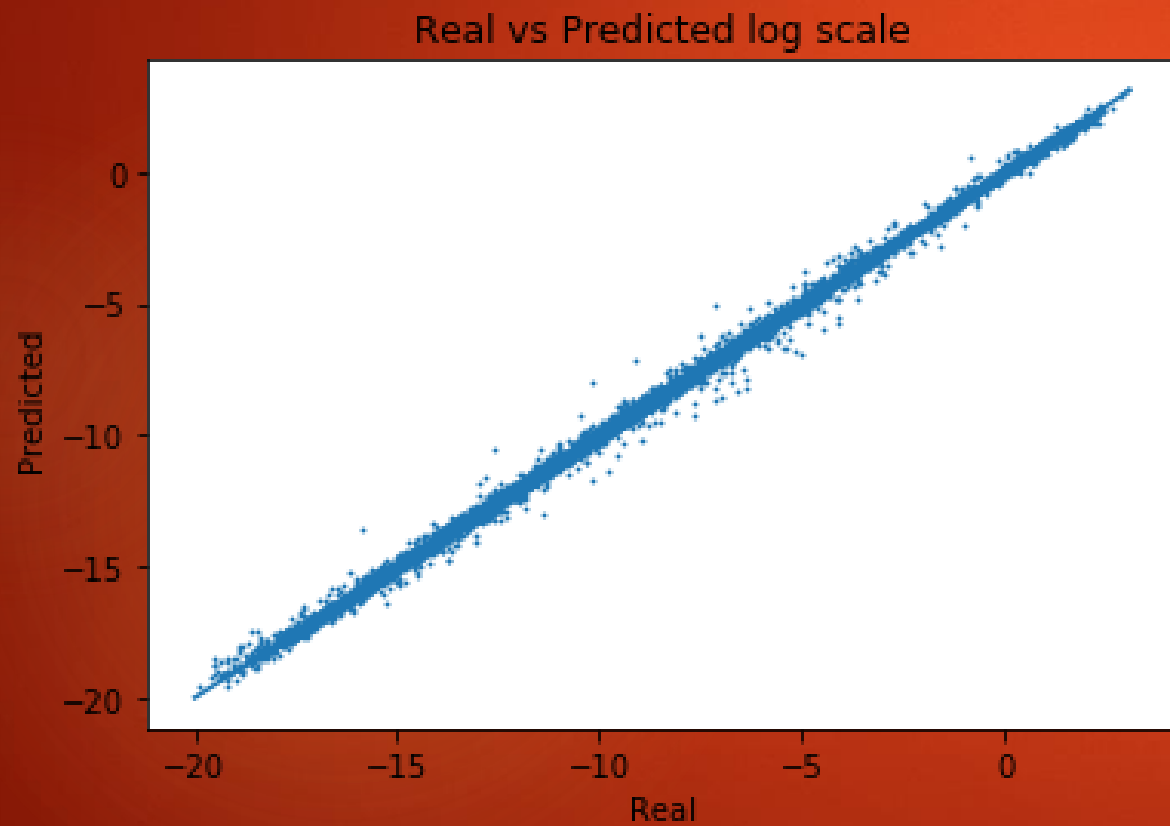
Percent Error



Summary

	Inside 1 SD	Inside 2 SDs	Inside 3 SD3	Outside of 3 SDs	Below min	Above max	Percent Error
Flipout Batched PRELU	25.59	48.12	65.86	34.17	9.01	22.04	11.40
Reparameterization Batched PRELU	39.31	67.16	83.00	17.00	5.02	9.65	13.21
Normal Dense PRELU	N/A	N/A	N/A	N/A	N/A	N/A	18.35
HMC Relu 1000	46.55	74.33	86.78	13.22	7.33	5.70	9.79
HMC Relu 5000	55.95	81.65	91.29	8.71	4.41	2.90	9.01
HMC Relu 195 from 3900	64.61	86.97	94.08	5.92	4.64	3.16	8.11

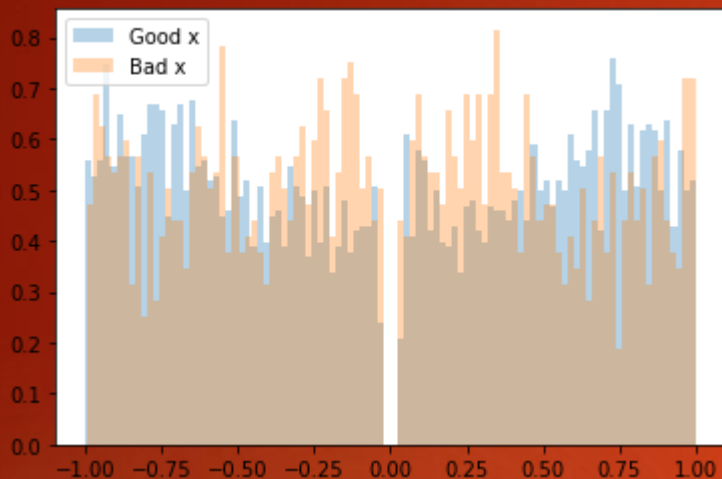
Real vs. Predicted



Error Analysis

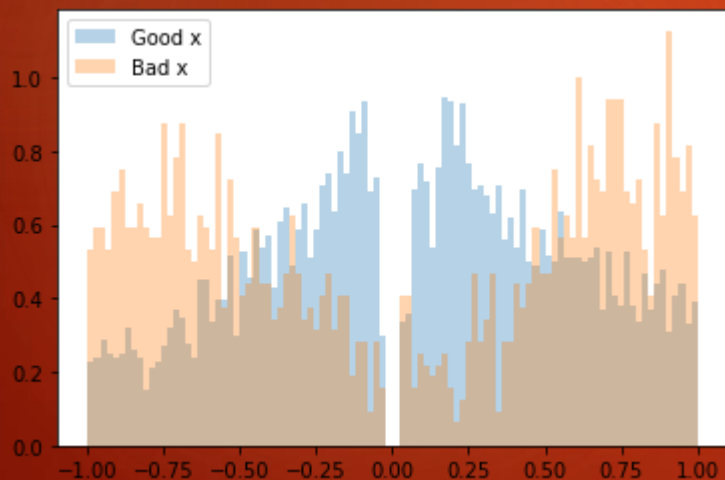
Wino mass parameter

x 1



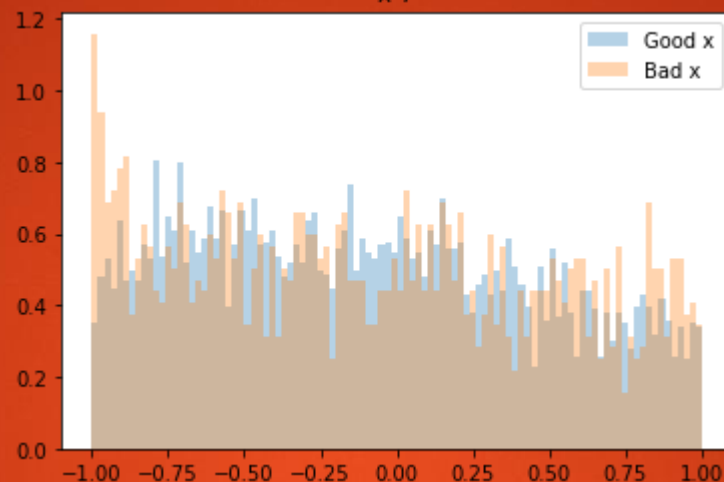
Higgsino mass parameter

x 6



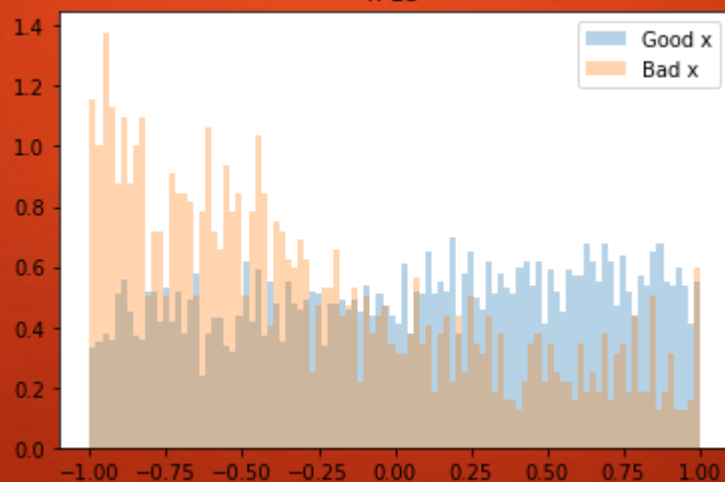
Ratio of the vacuum
expectation values of
the two Higgs doublets

x 7



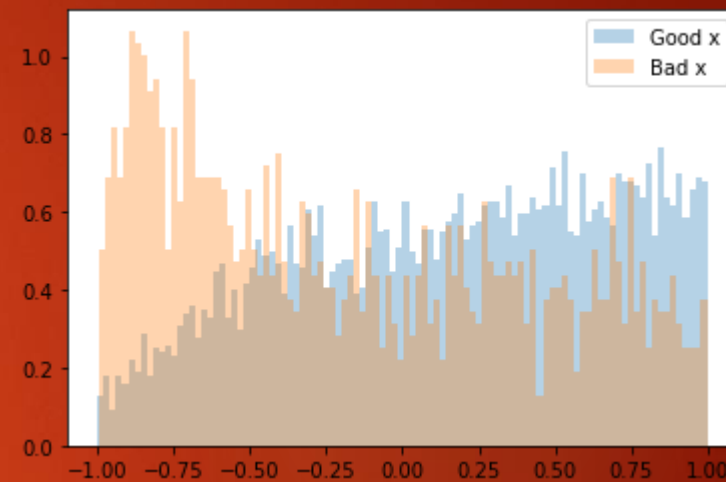
m_q

x 13



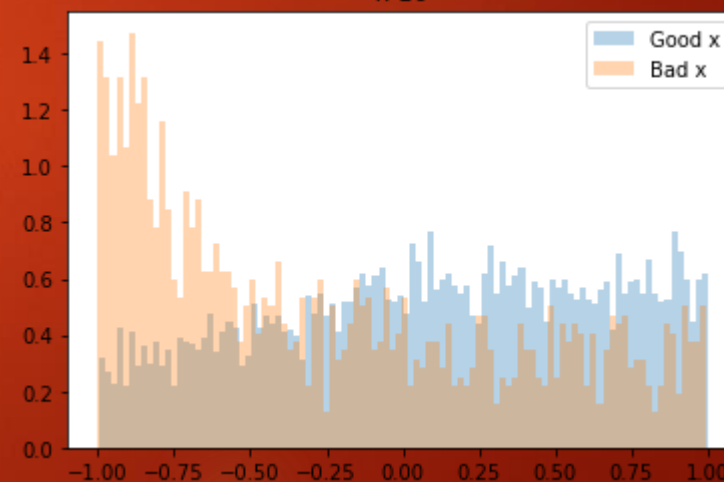
m_Q

x 14



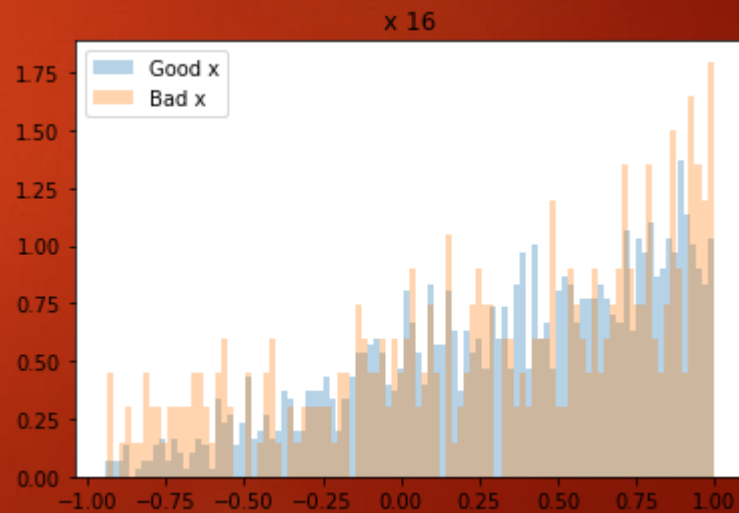
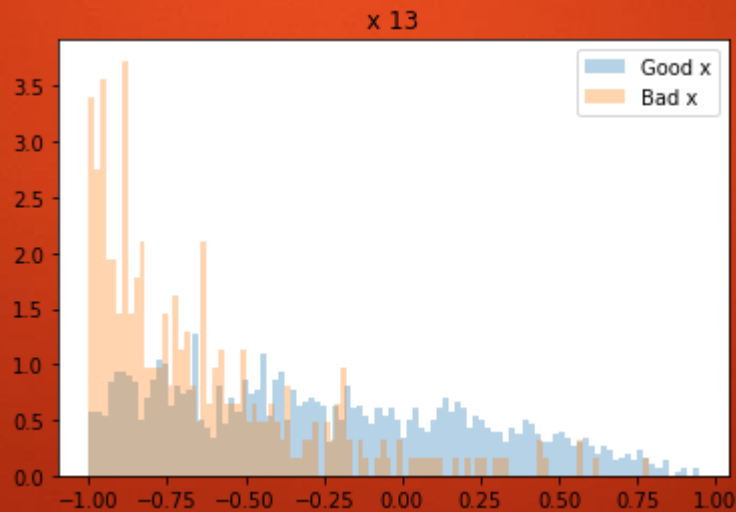
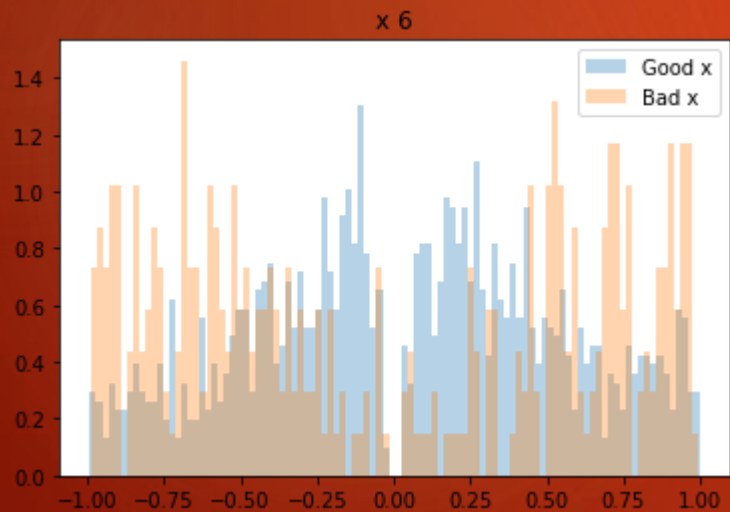
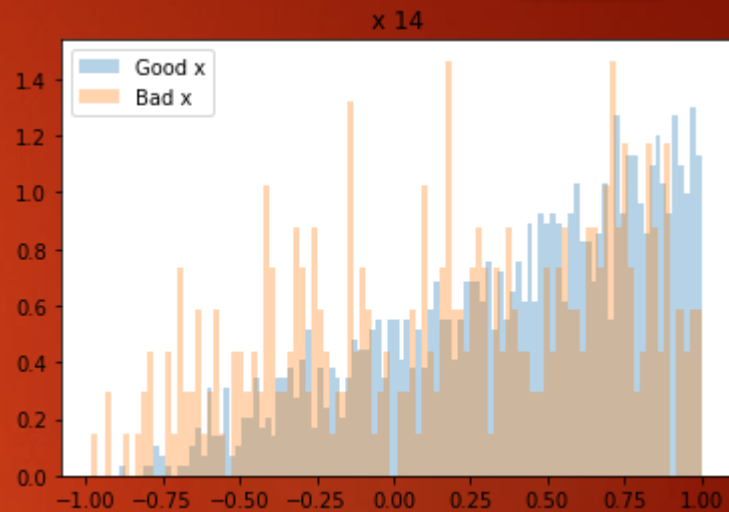
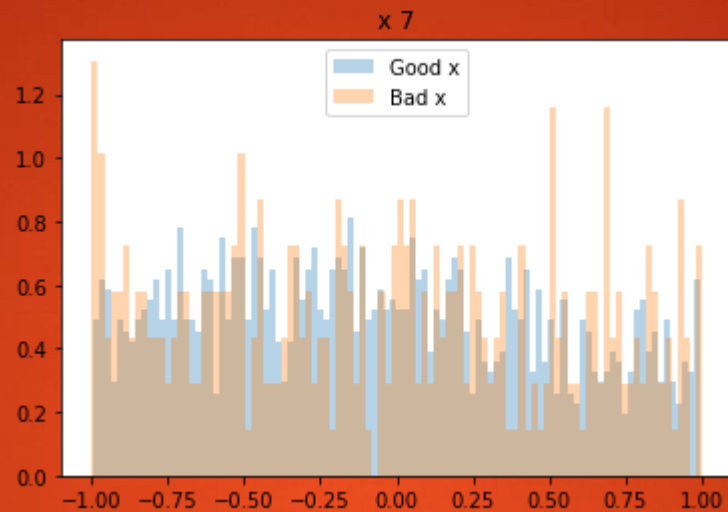
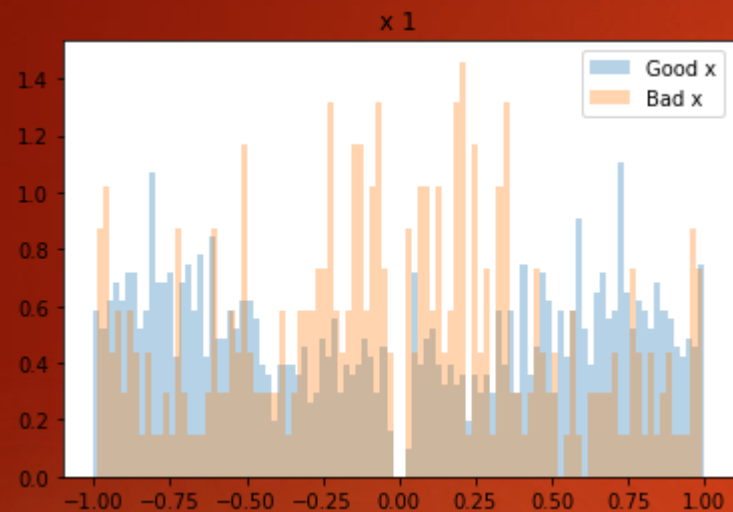
Stop quark mass parameter

x 16



$$m_q < m_Q \text{ and } m_u < m_t$$

5,185 points
6.9 % Error



Goals for next week

- ▶ Finish poster
- ▶ Research parameter relationships and apply this to data analysis
- ▶ Fix non invertible error